

## SEQUENCE LISTING

&lt;110&gt; NORRIS, STEVEN J.

&lt;120&gt; VMP-LIKE SEQUENCES OF PATHOGENIC BORRELIA SPECIES AND STRAINS

&lt;130&gt; UTSH:264US

&lt;140&gt; UNKNOWN

&lt;141&gt; 2005-06-17

&lt;150&gt; PCT/US2003/04118

&lt;151&gt; 2003-12-22

&lt;150&gt; 60/435,077

&lt;151&gt; 2002-12-20

&lt;160&gt; 81

&lt;170&gt; PatentIn Ver. 2.1

&lt;210&gt; 1

&lt;211&gt; 1227

&lt;212&gt; DNA

&lt;213&gt; Borrelia burgdorferi

&lt;220&gt;

&lt;221&gt; CDS

&lt;222&gt; (75)..(1142)

&lt;400&gt; 1

acctacactt gttaaaactc tcttttttgag ttaagatgat aacttatact tttcattata 60

aggagacgat gaat atg aaa aaa att tca agt gca agt tta tta aca act 110

Met Lys Lys Ile Ser Ser Ala Ser Leu Leu Thr Thr

1 5 10

ttc ttt gtt ttt att aat tgt aaa agc caa gtt gct gat aag gac gac 158

Phe Phe Val Phe Ile Asn Cys Lys Ser Gln Val Ala Asp Lys Asp Asp

15 20 25

cca aca aac aaa ttt tac caa tct gtc ata caa tta ggt aac gga ttt 206

Pro Thr Asn Lys Phe Tyr Gln Ser Val Ile Gln Leu Gly Asn Gly Phe

30 35 40

ctt gat gta ttc aca tct ttt ggt ggg tta gta gca gag gct ttt gga 254

Leu Asp Val Phe Thr Ser Phe Gly Gly Leu Val Ala Glu Ala Phe Gly

45 50 55 60

ttt aaa tca gat cca aaa aaa tct gat gta aaa acc tat ttt act act 302

Phe Lys Ser Asp Pro Lys Lys Ser Asp Val Lys Thr Tyr Phe Thr Thr

65 70 75

gta gct gcc aaa ttg gaa aaa aca aaa acc gat ctt aat agt ttg cct 350

Val Ala Ala Lys Leu Glu Lys Thr Lys Thr Asp Leu Asn Ser Leu Pro

80 85 90

aag gaa aaa agc gat ata agt agt acg acg ggg aaa cca gat agt aca	398
Lys Glu Lys Ser Asp Ile Ser Ser Thr Thr Gly Lys Pro Asp Ser Thr	
95 100 105	
ggt tct gtt gga act gcc gtt gag ggg gct att aag gaa gtt agc gag	446
Gly Ser Val Gly Thr Ala Val Glu Gly Ala Ile Lys Glu Val Ser Glu	
110 115 120	
ttg ttg gat aag ctg gta aaa gct gta aag aca gct gag ggg gct tca	494
Leu Leu Asp Lys Leu Val Lys Ala Val Lys Thr Ala Glu Gly Ala Ser	
125 130 135 140	
agt ggt act gct gca att gga gaa gtt gtg gct gat gct gat gct gca	542
Ser Gly Thr Ala Ala Ile Gly Glu Val Val Ala Asp Ala Asp Ala Ala	
145 150 155	
aag gtt gct gat aag gcg agt gtg aag ggg att gct aag ggg ata aag	590
Lys Val Ala Asp Lys Ala Ser Val Lys Gly Ile Ala Lys Gly Ile Lys	
160 165 170	
gag att gtt gaa gct gct ggg ggg agt gaa aag ctg aaa gct gtt gct	638
Glu Ile Val Glu Ala Ala Gly Gly Ser Glu Lys Leu Lys Ala Val Ala	
175 180 185	
gct gct aaa ggg gag aat aat aaa ggg gca ggg aag ttg ttt ggg aag	686
Ala Ala Lys Gly Glu Asn Asn Lys Gly Ala Gly Lys Leu Phe Gly Lys	
190 195 200	
gct ggt gct gct gct cat ggg gac agt gag gct gct agc aag gcg gct	734
Ala Gly Ala Ala Ala His Gly Asp Ser Glu Ala Ala Ser Lys Ala Ala	
205 210 215 220	
ggt gct gtt agt gct gtt agt ggg gag cag ata tta agt gcg att gtt	782
Gly Ala Val Ser Ala Val Ser Gly Glu Gln Ile Leu Ser Ala Ile Val	
225 230 235	
acg gct gct gat gcg gct gag cag gat gga aag aag cct gag gag gct	830
Thr Ala Ala Asp Ala Ala Glu Gln Asp Gly Lys Lys Pro Glu Glu Ala	
240 245 250	
aaa aat ccg att gct gct gct att ggg gat aaa gat ggg ggt gcg gag	878
Lys Asn Pro Ile Ala Ala Ala Ile Gly Asp Lys Asp Gly Gly Ala Glu	
255 260 265	
ttt ggt cag gat gag atg aag aag gat gat cag att gct gct gct att	926
Phe Gly Gln Asp Glu Met Lys Lys Asp Asp Gln Ile Ala Ala Ala Ile	
270 275 280	
gct ttg agg ggg atg gct aag gat gga aag ttt gct gtg aag gat ggt	974
Ala Leu Arg Gly Met Ala Lys Asp Gly Lys Phe Ala Val Lys Asp Gly	
285 290 295 300	
gag aaa gag aag gct gag ggg gct att aag gga gct gct gag tct gca	1022
Glu Lys Glu Lys Ala Glu Gly Ala Ile Lys Gly Ala Ala Glu Ser Ala	
305 310 315	
gtt cgc aaa gtt tta ggg gct att act ggg cta ata gga gac gcc gtg	1070

Val Arg Lys Val Leu Gly Ala Ile Thr Gly Leu Ile Gly Asp Ala Val  
320 325 330

agt tcc ggg cta agg aaa gtc ggt gat tca gtg aag gct gct agt aaa 1118  
Ser Ser Gly Leu Arg Lys Val Gly Asp Ser Val Lys Ala Ala Ser Lys  
335 340 345

gaa aca cct cct gcc ttg aat aag tgatttaatt aagtgtatgg acacgactat 1172  
Glu Thr Pro Pro Ala Leu Asn Lys  
350 355

gccctcatga ttgaggaaat agtcgagaga tatatatatact aaaagataat aaata 1227

<210> 2  
<211> 356  
<212> PRT  
<213> Borrelia burgdorferi

<400> 2  
Met Lys Lys Ile Ser Ser Ala Ser Leu Leu Thr Thr Phe Phe Val Phe  
1 5 10 15

Ile Asn Cys Lys Ser Gln Val Ala Asp Lys Asp Asp Pro Thr Asn Lys  
20 25 30

Phe Tyr Gln Ser Val Ile Gln Leu Gly Asn Gly Phe Leu Asp Val Phe  
35 40 45

Thr Ser Phe Gly Gly Leu Val Ala Glu Ala Phe Gly Phe Lys Ser Asp  
50 55 60

Pro Lys Lys Ser Asp Val Lys Thr Tyr Phe Thr Thr Val Ala Ala Lys  
65 70 75 80

Leu Glu Lys Thr Lys Thr Asp Leu Asn Ser Leu Pro Lys Glu Lys Ser  
85 90 95

Asp Ile Ser Ser Thr Thr Gly Lys Pro Asp Ser Thr Gly Ser Val Gly  
100 105 110

Thr Ala Val Glu Gly Ala Ile Lys Glu Val Ser Glu Leu Leu Asp Lys  
115 120 125

Leu Val Lys Ala Val Lys Thr Ala Glu Gly Ala Ser Ser Gly Thr Ala  
130 135 140

Ala Ile Gly Glu Val Val Ala Asp Ala Asp Ala Ala Lys Val Ala Asp  
145 150 155 160

Lys Ala Ser Val Lys Gly Ile Ala Lys Gly Ile Lys Glu Ile Val Glu  
165 170 175

Ala Ala Gly Gly Ser Glu Lys Leu Lys Ala Val Ala Ala Ala Lys Gly  
180 185 190

Glu Asn Asn Lys Gly Ala Gly Lys Leu Phe Gly Lys Ala Gly Ala Ala

195	200	205
Ala His Gly Asp Ser Glu	Ala Ala Ser Lys	Ala Ala Gly Ala Val Ser
210	215	220
Ala Val Ser Gly Glu Gln	Ile Leu Ser Ala	Ile Val Thr Ala Ala Asp
225	230	235 240
Ala Ala Glu Gln Asp Gly	Lys Lys Pro Glu	Glu Ala Lys Asn Pro Ile
245	250	255
Ala Ala Ala Ile Gly Asp	Lys Asp Gly Gly	Ala Glu Phe Gly Gln Asp
260	265	270
Glu Met Lys Lys Asp Asp	Gln Ile Ala Ala	Ile Ala Leu Arg Gly
275	280	285
Met Ala Lys Asp Gly Lys	Phe Ala Val Lys Asp	Gly Glu Lys Glu Lys
290	295	300
Ala Glu Gly Ala Ile Lys	Gly Ala Ala Glu Ser	Ala Val Arg Lys Val
305	310	315 320
Leu Gly Ala Ile Thr Gly	Leu Ile Gly Asp	Ala Val Ser Ser Gly Leu
325	330	335
Arg Lys Val Gly Asp Ser	Val Lys Ala Ala	Ser Lys Glu Thr Pro Pro
340	345	350
Ala Leu Asn Lys		
355		

<210> 3  
 <211> 1141  
 <212> DNA  
 <213> Borrelia hermsii

<220>  
 <221> CDS  
 <222> (1)..(1062)

<400> 3																
atg	aga	aaa	aga	ata	agt	gca	ata	ata	atg	act	tta	ttt	atg	gta	tta	48
Met	Arg	Lys	Arg	Ile	Ser	Ala	Ile	Ile	Met	Thr	Leu	Phe	Met	Val	Leu	
1				5					10					15		
gta agc tgt aat agc ggt ggg gtt gcg gaa gat cct aaa act gtg tat															96	
Val	Ser	Cys	Asn	Ser	Gly	Gly	Val	Ala	Glu	Asp	Pro	Lys	Thr	Val	Tyr	
			20					25					30			
tta aca tct ata gct aat tta ggg aaa gga ttt tta gat gtt ttt gtg															144	
Leu	Thr	Ser	Ile	Ala	Asn	Leu	Gly	Lys	Gly	Phe	Leu	Asp	Val	Phe	Val	
			35				40					45				
act ttt gga gat atg gtt act gga gct ttt ggt att aag gca gat act															192	

Thr	Phe	Gly	Asp	Met	Val	Thr	Gly	Ala	Phe	Gly	Ile	Lys	Ala	Asp	Thr		
50						55					60						
aag	aaa	agt	gat	ata	ggg	aag	tat	ttt	act	gat	att	gag	agc	act	atg	240	
Lys	Lys	Ser	Asp	Ile	Gly	Lys	Tyr	Phe	Thr	Asp	Ile	Glu	Ser	Thr	Met		
65					70					75					80		
aca	tca	gtt	aaa	aag	aag	ttg	caa	gat	gaa	gtt	gct	aag	aat	ggg	aac	288	
Thr	Ser	Val	Lys	Lys	Lys	Leu	Gln	Asp	Glu	Val	Ala	Lys	Asn	Gly	Asn		
				85					90					95			
tat	cca	aag	gta	aag	aca	gct	gtt	gac	gaa	ttt	gtt	gca	atc	tta	gga	336	
Tyr	Pro	Lys	Val	Lys	Thr	Ala	Val	Asp	Glu	Phe	Val	Ala	Ile	Leu	Gly		
			100					105					110				
aag	atc	gag	aaa	gga	gca	aaa	gaa	gca	tct	aaa	ggg	gct	act	ggg	gat	384	
Lys	Ile	Glu	Lys	Gly	Ala	Lys	Glu	Ala	Ser	Lys	Gly	Ala	Thr	Gly	Asp		
		115					120					125					
gtt	att	att	ggg	aat	act	gtt	aag	aat	ggg	gat	gct	gta	cct	gga	gaa	432	
Val	Ile	Ile	Gly	Asn	Thr	Val	Lys	Asn	Gly	Asp	Ala	Val	Pro	Gly	Glu		
		130				135					140						
gca	aca	agt	gtc	aat	tct	ctt	gtt	aaa	gga	att	aaa	gaa	ata	gtt	ggg	480	
Ala	Thr	Ser	Val	Asn	Ser	Leu	Val	Lys	Gly	Ile	Lys	Glu	Ile	Val	Gly		
145					150					155					160		
gta	gtc	ttg	aag	gaa	ggg	aag	gca	gat	gct	gat	gct	act	aaa	gat	gat	528	
Val	Val	Leu	Lys	Glu	Gly	Lys	Ala	Asp	Ala	Asp	Ala	Thr	Lys	Asp	Asp		
			165						170					175			
agt	aag	aaa	gat	att	ggg	aaa	tta	ttt	acc	gca	acc	act	gat	gcg	aat	576	
Ser	Lys	Lys	Asp	Ile	Gly	Lys	Leu	Phe	Thr	Ala	Thr	Thr	Asp	Ala	Asn		
			180					185					190				
aga	gct	gat	aat	gcg	gca	gct	caa	gca	gct	gca	gcg	tca	ata	gga	gca	624	
Arg	Ala	Asp	Asn	Ala	Ala	Ala	Gln	Ala	Ala	Ala	Ala	Ser	Ile	Gly	Ala		
			195				200					205					
gtg	aca	ggg	gct	gat	atc	ttg	caa	gct	ata	gta	caa	tct	aag	gaa	aat	672	
Val	Thr	Gly	Ala	Asp	Ile	Leu	Gln	Ala	Ile	Val	Gln	Ser	Lys	Glu	Asn		
		210				215					220						
cct	gtt	gca	aat	agt	act	gat	gga	att	gaa	aaa	gca	aca	gat	gca	gct	720	
Pro	Val	Ala	Asn	Ser	Thr	Asp	Gly	Ile	Glu	Lys	Ala	Thr	Asp	Ala	Ala		
225					230					235				240			
gag	att	gca	gtt	gct	cca	gct	aaa	gat	aat	aaa	aaa	gag	att	aaa	gat	768	
Glu	Ile	Ala	Val	Ala	Pro	Ala	Lys	Asp	Asn	Lys	Lys	Glu	Ile	Lys	Asp		
				245					250					255			
gga	gca	aaa	aaa	gac	gca	gtt	att	gct	gca	ggc	att	gca	ctg	cga	gca	816	
Gly	Ala	Lys	Lys	Asp	Ala	Val	Ile	Ala	Ala	Gly	Ile	Ala	Leu	Arg	Ala		
			260					265					270				
atg	gct	aag	aat	ggg	aca	ttt	tct	att	aaa	aac	aat	gaa	gat	gcg	gct	864	
Met	Ala	Lys	Asn	Gly	Thr	Phe	Ser	Ile	Lys	Asn	Asn	Glu	Asp	Ala	Ala		

275	280	285	
gta acg acg ata aat agt gca gca gca agc gca gtg aac aag att tta			912
Val Thr Thr Ile Asn Ser Ala Ala Ala Ser Ala Val Asn Lys Ile Leu			
290	295	300	
agc act cta ata ata gca ata agg aat aca gtt gat agt ggt tta aaa			960
Ser Thr Leu Ile Ile Ala Ile Arg Asn Thr Val Asp Ser Gly Leu Lys			
305	310	315	320
aca ata aat gag gct ctt gct aca gtt aaa caa gaa gat aaa tct gta			1008
Thr Ile Asn Glu Ala Leu Ala Thr Val Lys Gln Glu Asp Lys Ser Val			
325	330	335	
gaa gca act aat act gca gaa gca aca act agt ggt cag caa gcg aaa			1056
Glu Ala Thr Asn Thr Ala Glu Ala Thr Thr Ser Gly Gln Gln Ala Lys			
340	345	350	
aac tag ttaagggtaa atataaagga taaagttatt gtaagggaaa agcttttctt			1112
Asn			
gtttttaatg caggaatgta gtttctctg			1141

<210> 4

<211> 353

<212> PRT

<213> *Borrelia hermsii*

<400> 4

Met Arg Lys Arg Ile Ser Ala Ile Ile Met Thr Leu Phe Met Val Leu			
1	5	10	15
Val Ser Cys Asn Ser Gly Gly Val Ala Glu Asp Pro Lys Thr Val Tyr			
20	25	30	
Leu Thr Ser Ile Ala Asn Leu Gly Lys Gly Phe Leu Asp Val Phe Val			
35	40	45	
Thr Phe Gly Asp Met Val Thr Gly Ala Phe Gly Ile Lys Ala Asp Thr			
50	55	60	
Lys Lys Ser Asp Ile Gly Lys Tyr Phe Thr Asp Ile Glu Ser Thr Met			
65	70	75	80
Thr Ser Val Lys Lys Lys Leu Gln Asp Glu Val Ala Lys Asn Gly Asn			
85	90	95	
Tyr Pro Lys Val Lys Thr Ala Val Asp Glu Phe Val Ala Ile Leu Gly			
100	105	110	
Lys Ile Glu Lys Gly Ala Lys Glu Ala Ser Lys Gly Ala Thr Gly Asp			
115	120	125	
Val Ile Ile Gly Asn Thr Val Lys Asn Gly Asp Ala Val Pro Gly Glu			
130	135	140	
Ala Thr Ser Val Asn Ser Leu Val Lys Gly Ile Lys Glu Ile Val Gly			
145	150	155	160
Val Val Leu Lys Glu Gly Lys Ala Asp Ala Asp Ala Thr Lys Asp Asp			
165	170	175	
Ser Lys Lys Asp Ile Gly Lys Leu Phe Thr Ala Thr Thr Asp Ala Asn			
180	185	190	
Arg Ala Asp Asn Ala Ala Ala Gln Ala Ala Ala Ala Ser Ile Gly Ala			
195	200	205	
Val Thr Gly Ala Asp Ile Leu Gln Ala Ile Val Gln Ser Lys Glu Asn			

210	215	220
Pro Val Ala Asn Ser Thr Asp Gly Ile Glu Lys Ala Thr Asp Ala Ala		
225	230	235
Glu Ile Ala Val Ala Pro Ala Lys Asp Asn Lys Lys Glu Ile Lys Asp		240
	245	250
Gly Ala Lys Lys Asp Ala Val Ile Ala Ala Gly Ile Ala Leu Arg Ala		255
	260	265
Met Ala Lys Asn Gly Thr Phe Ser Ile Lys Asn Asn Glu Asp Ala Ala		270
	275	280
Val Thr Thr Ile Asn Ser Ala Ala Ser Ala Val Asn Lys Ile Leu		285
	290	295
Ser Thr Leu Ile Ile Ala Ile Arg Asn Thr Val Asp Ser Gly Leu Lys		300
305	310	315
Thr Ile Asn Glu Ala Leu Ala Thr Val Lys Gln Glu Asp Lys Ser Val		320
	325	330
Glu Ala Thr Asn Thr Ala Glu Ala Thr Thr Ser Gly Gln Gln Ala Lys		335
	340	345
		350

Asn

<210> 5  
 <211> 8762  
 <212> DNA  
 <213> Borrelia afzelii

<220>  
 <221> CDS  
 <222> (8226) .. (8762)

<400> 5  
 gagagtgctg ttgatggggt tagcaagtgg ttagaagaga tgataaaagc tgctaaggag 60  
 gctgctacaa aggggtggtac tgggtggtgg agcgaaaaga ttggggatgt tgggtgctgct 120  
 aataatcagg gtgctgtagc tgataaggac agtggttaagg ggattgcgaa ggggataaag 180  
 gggattgttg atgctgctgg gaaggctttt ggtaaggatg gtaatgcgct gacaggtgta 240  
 aaagaagttg ctgatgaggc tggggctaac gaggatgcgg ggaagttggt tgctggtaat 300  
 gctggtaatg ctgctgctgc tgacattgcg aaggcggctg gtgctgttac tgcggttagt 360  
 ggggagcaga tactgaaagc tattgttgat ggtgctggtg gtgcggctca agatggtaaa 420  
 aaggctgcgg aggctaagaa tccgattgca gctgcgattg gggctgatgc tgctggtgcg 480  
 gatttttggtg atgatatgaa gaagagtgat aagattgctg cggctattgt tttgaggggg 540  
 gtggctaaga gtggaaagtt tgctgttgct aatgctgcta agaaggagag tgtgaagagt 600  
 gctgtggaga gtgctgttga tgaggtttagc aagtggttag aagagatgat aaaagctgct 660  
 ggtggggctg ctaagggtgg tactggtggt aataacgaaa agattgggga ttctgataat 720  
 aataagggtg ctgtagctga taaggacagt gttaagggga ttgcgaaggg gataaagggg 780

attgttgatg ctgctgggaa ggcttttggg aaggatggta atgcgctgaa ggatgttgca 840  
aaagttgctg atgatgcggc tggggctaac gcgaatgcag ggaagttggt tgctggtaat 900  
gctgctggtg gtgccgctga tgctgatgat gctaacattg cgaaggcggc tggtgctggt 960  
agtgcggtta gtggggagca gatactgaaa gctattgttg atgctgctgg tgctgctgct 1020  
aatcaggatg gtaagaaggc tgcggatgct aagaatccga ttgcagctgc gattgggact 1080  
aatgatgatg gggcggagtt taaggatgga atgaagaaga gtgataatat tgctgcagct 1140  
attgttttga ggggggtggc taagggtgga aagtttgctg ttgctaatac tgctaataat 1200  
agtaaggcga gtgtgaagag tgctgtggag agtgctgttg atgaggttag caagtgggta 1260  
gaagagatga taacagctgc tggtgaggct gctacaaagg gtggtgatgc tgggtggtgg 1320  
gctgataaga ttggggatgt tggtgctgct aataatgggt ctgtagctga tgcgagcagt 1380  
gttaaggaga ttgcgaaggg gataaagggg attgttgatg ctgctgggaa ggcttttggc 1440  
aaggatggta atgcgctgaa ggatgttgca gaagttgctg atgataagaa ggaggcgggg 1500  
aagttgtttg ctggtaatgc tgggtggtgct gttgctgatg ctgctgcgat tgggaaggcg 1560  
gctggtgctg ttactgcggt tagtggggag cagatactga aagctattgt tgatgctgct 1620  
ggtggtgcgg atcaggcggg taagaaggct gatgcggcta agaatccgat tgcagctgcg 1680  
attggggctg atgctgctgg tgctggtgcg gatttttgga atgatatgaa gaagagaaat 1740  
gataagattg ttgcggctat tgttttgagg ggggtggcta aggatggaaa gtttgctgct 1800  
gctgctaata atgataatag taaggcgagt gtgaagagt ctgtggagag tgctgttgat 1860  
gaggttagca agtggttaga agagatgata acagctgctg atggggctgc taaagggtgg 1920  
actggtggta atagcgaata gattggggat gctggtgata ataataatgg tgctgtagct 1980  
gatgagaaca gtgttaagga gattgcaaag gggataaagg ggattgttgc ggctgctggg 2040  
aaggcttttg gcaaggatgg caaggatggg gatgcgctga aggatgttga aacagttgct 2100  
gctgagaatg aggctaaca ggatgcgggg aagttgtttg ctggtgctaa tggtaatgct 2160  
ggtgctgctg ttggtgacat tgcgaaggcg gctgctgctg ttactgcggt tagtggggag 2220  
cagatactaa aagctattgt tgatgctgct ggtgatgcgg atcaggcggg taagaaggct 2280  
gctgaggcta agaatccgat tgcagctgcg attggggcta atgctgctga taatgcggcg 2340  
gcgtttggta aggatgagat gaagaagagt gataagattg ctgcagctat tgttttgagg 2400  
ggggtggcta aggatggaaa gtttgctggt gctaatagcta atgatgataa gaaggcgagt 2460  
gtgaagagt ctgtggagag tgctgtggat gaggttagca agtggttaga agagatgata 2520



acagctgcta aggaggctgc tacaaagggg ggtactgggtg gtaataacga aaagattgga 2580  
gattctgatg ctaataatgg tgcgaaggct gatgcgagca gtgttaatgg gattgcgaat 2640  
gggataaagg ggattgttga tgctgctggg aaggcttttg gcaaggaggg tagtgcgctg 2700  
aaggatgtta aaacagttgc tgctgagaat gaggctaaca aggatgcggg gaagttgttt 2760  
gctggtaaga atggtaatgc tgatgctgct gatgctgctg acattgcgaa ggcggctggg 2820  
gctgttagtg cggtagtggt ggagcagata ctgaaagcta ttgttgatgg tgctgggtgat 2880  
gcagctaatc aggcgggtaa aaaggctgct gaggctaaga atccgattgc ggctgcgatt 2940  
gggactaatg aagctggggc ggagtttggt gatgatatga agaagagaaa tgataagatt 3000  
gctgcggcta ttgttttgag ggggggtggc aaggatggaa agtttgctgt tgctaattgct 3060  
gctgctgata atagtaaggc gagtgtgaag agtgctgttg atgaggttag caagtgggta 3120  
gaagagatga taaaggctgc tggtagggc gctacaaagg gtggatgatgc tggtaggtgg 3180  
gctgataaga ttggggatgc tggtagaag ggtgctgtag ctgatgcgag cagtgttaag 3240  
gagattgcga atgggataaa ggggattgtt gatgctgctg ggaaggcttt tggcaaggag 3300  
ggtagtgcgc tgaaggatgt taaaacagtt gctgctgaga atgaggctaa caaggatgcg 3360  
gggaagttgt ttgctggtaa tgctggtaat ggtgctgctg atgacattgc gaaggcggct 3420  
gctgctgtta ctgcgggttag tggggagcag atactgaaag ctattgttga tgctgctggg 3480  
gataaggcta atcaggatgg taaaaaggct gcggatgcta agaattccgat tgcggctgcg 3540  
attggggctg ctgatgctgg tgctgcggcg gcgtttaatg agaattgat gaagaagagt 3600  
gataagattg ctgcagctat tgttttgagg ggggtggcta aggatggaaa gtttgctgct 3660  
gctgatgctg atgctaataa tagtaaggcg agcgtgaaga gtgctgttgg tgaggttagc 3720  
aagtggttag aagagatgat aaaagctgct ggtgaggctg caaaagttgg tggtaggtg 3780  
ggtagcgaag agattgggga tgctgataat aataagggtg ctgtagctga tgcgagcagt 3840  
gttaatggga ttgcgaatgg gataaagggg attgttgatg ctgctgggaa ggcttttggt 3900  
aaggatgggt cgctggcagg tgttgagct gctgctgaga atgatgataa gaaggatgcg 3960  
gggaagttgt ttgctggtaa gaatgggtgg gctgggtgctg ctgatgcgat tgggaaggcg 4020  
gctgctgctg ttactgcggt tagtggggag cagatactga aagctattgt tgatgctgct 4080  
ggtgctgcag ctaatcaggc gggtaaaaag gctgcggatg ctaagaatcc gattgcggct 4140  
gcgattggga ctgctgatga tggggcggag tttaaggatg atatgaagaa gagtgataat 4200

attgctgcgg ctattgtttt gaggggggtg gctaaggatg gaaagtttgc tgttgctaata 4260  
gctgatgata ataaggcgag tgtgaagagt gctgtggaga gtgctgttga tgaggtagc 4320  
aagtggtag aagagatgat aacagctgct ggtgaggctg caaaagttgg tgctggtagt 4380  
ggtgctgata agattgggga tgctgctaata aatcagggtg cgaaggctga tgagagcagt 4440  
gttaatggaa ttgcaaagg gataaagggtt attgttgatg ctgctgggaa ggcttttggc 4500  
aaggagggtg gtgcgctgaa ggatgttgca aaagttgctg atgatgataa caaggatgcg 4560  
gggaagttgt ttgctggtaa tgctggtagt ggtgctggtag ctgatattgc gaaggcggct 4620  
gctgctgtta ctgcggttag tggggagcag atactgaaag ctattgttga tgctgctggt 4680  
gctgcggatc aggcgggtgc agctgctggt gcggctaaga atccgattgc ggctgcgatt 4740  
ggggctgatg ctggtgctgc ggaggagttt aaggatgaga tgaagaagag tgataagatt 4800  
gctgcggcta ttgttttgag gggggtggct aagggtggaa agtttgctgt tgctgctaata 4860  
gatgctgcaa atgtgaagag tgctgtggag agtgctgttg gtgaggtag cgcattggtta 4920  
gaagagatga taacagctgc tagtgaggct gctacaaagg gtggtactgg tggtagtgg 4980  
ggtgatagtg aaaagattgg ggattctgat gctaataatg gtgctgtagc tgatgcgagc 5040  
agtgttaagg agattgcgaa ggggataaag gggattgttg atgctgctgg gaaggctttt 5100  
ggtaaggatg gtaatgcgct gaaggatgtt gcagaagttg ctgatgatga ggctaacgcg 5160  
gatgcgggga agttgtttgc tggtaatgct ggtaatgctg ctgctgctga cgttgcaag 5220  
gcggctggtg ctgttactgc ggtagtggg gagcagatac tgaaagctat tgttgatgct 5280  
gctggtgctg cggatcaggc gggtgcaaag gctgatgcgg ctaagaatcc gattgcagct 5340  
gcgattggga ctaatgaagc tggggcgcg tttaaggatg gaatgaagaa gagaaatgat 5400  
aatattgctg cggctattgt tttgaggggg gtggctaaga gtggaaagt ttgctgttgc 5460  
gctgctgatg ctggtaaggc gagagtgtga agagtgtgt ggagagtgt gttgatgagg 5520  
ttagcaagtg gttagaagag atgataacag ctgctagtga ggctgcaaaa gttggtgctg 5580  
gtggtgatga taagattggg gattctgcta ataatggtgc ttagctgat gcgggcagt 5640  
ttaagggaat tgcgaagggt ataaagggtt ttgttgatgc tgctgggaag gcttttggta 5700  
aggagggtga tgcgctgaag gatgttgcaa aagttgctga tgagaatggg gataacaagg 5760  
atgcggggaa gttgtttgct ggtgagaatg gtaatgctgg tgggtgctgct gatgctgaca 5820  
ttgcgaaggc ggctgctgct gttactgcgg ttagtgggga gcagatactg aaagctattg 5880  
ttgaggctgc tggtagtggg gatgcagcta atcaggcggg taagaaggct gatgaggcta 5940

agaatccgat tgcggctgctg attgggactg atgatgctgg ggcggcgctt ggtcaggatg 6000  
atatgaagaa gagaaatgat aatattgctg cggctattgt tttgaggggg gtggctaagg 6060  
gtggaaagtt tgctgttgct aatgctgcta atgatatga ggcgagtgtg aagagtgtgct 6120  
tggagagtgc tggtgatgag gtttagcaagt ggtagaaga gataataaca gctactggga 6180  
aggcttttgg taaggatggt aatgcgctgg cagggtgtgc aaaagttgct gatgatgagg 6240  
ctaacgcgga tgcggggaag ttgtttgctg gtgagaatgg taatgctggt gctgctgcga 6300  
ttgggaaggc ggctgctgct gttactgcgg ttagtgggga gcagatactg aaagctattg 6360  
ttgatgctgc tgggtggtgc gctcagggtg gtgctggtgc tgggtcggct acgaatccga 6420  
ttgcagctgc gattggggct gctggtgatg gtgcggattt tggtaaggat gagatgaaga 6480  
agagaaatga taagattgct gcggctattg ttttgagggg ggtggctaag gatggaaagt 6540  
ttgctgctgc tgctaattgat agtaaggcga gtgtgaagag tgctgtggag agtgcgtgtg 6600  
atgaggtag caagtggta gaagagatga taacagctgc tgatgctgct gctgctaaag 6660  
ttggcgatgc tgggtggtggt gctgataaga ttggggatgt tgggtgctgct aataaggggtg 6720  
cgaaggctga tgcgagcagt gttaaggaga ttgcgaaggg gataaagggg attgttgatg 6780  
ctgctgggaa ggcttttggg ggtgatgctg tgaaggatgt taaagctgct ggtgatgata 6840  
acaaggaggc agggaaagttg tttgctggtg ctaatggtaa tgctggtgct aatgctgctg 6900  
ctgctgatga cattgcgaag gcggctggtg ctgttagtgc ggttagtggg gagcagatac 6960  
tgaaagctat tggtgaggcg gctggtgctg cggatcaggc ggggtgtaaag gctgaggagg 7020  
ctaagaatcc gattgcagct gcgattggga ctgatgatgc tgggtgcggcg gagtttggtg 7080  
agaatgatat gaagaagaat gataatattg ctgcggctat tgttttgagg ggggtggcta 7140  
agagtggaaa gtttgctgct aatgctaatt atgctggtaa gaaggagagt gtgaagagtg 7200  
ctgtggatga ggctagcaag tggtagaag agatgataac agctgctggt gaggctgcta 7260  
caaaggggtg tactggtgaa gctagcgaag agattgggga tggtggtgat aataatcatg 7320  
gtgctgtagc tgatgcggac agtgtaagg ggattgcga ggggataaag gggattgttg 7380  
atgctgctgg gaaggctttt ggtaaggatg gtgcgctgaa ggatgttgca gctgctgctg 7440  
gtgatgaggc taacaaggat gcggggaagt tgtttgctgg tcaggatggt ggtggtgctg 7500  
atggtgacat tgcgaaggcg gctgctgctg ttactgcggt tagtggggag cagatactga 7560  
aagctattgt tgaggctgct ggtgataagg ctaatcaggt ggggtgtaaag gctgctggtg 7620

cggctacgaa tccgattgca gctgcgattg ggactgatga tgataatgcg gcggcgtttg 7680  
 ataaggatga gatgaagaag agtaatgata agattgctgc ggctattggtt ttgagggggg 7740  
 tggctaagga tggaaaagttt gctgctaattg ctaatgataa tagtaaggcg agtgtgaaga 7800  
 gtgctgtgga tgaggtttagc aagtggtttag aagagatgat aacagctgct agtgatgctg 7860  
 ctacaaaaggg tggtagtggg gaagctagcg aaaagattgg ggattctgat gctaataagg 7920  
 gtgctggtgc tggggcgggcg tttggtgaga atgatatgaa gaagagaaat gataatattg 7980  
 ctgcagctat tgttttgagg ggggtggcta aggatggaaa gtttgctggtt aaggaggatt 8040  
 attgaactca gctttatagg ggaacagcaa ttcgctagaa aatgattaaa aagcttaact 8100  
 tcgactgggtt cttgccttaa ttttattcct ttgttattat ttatcaatta aattcacttc 8160  
 ggtttgcttt taaattaatt ctggtatact atgtatacta gatacacaaa ttaaggagaa 8220  
 gtgaa atg gaa aaa ata gaa aaa ttt aaa aac aaa tgt caa cat aaa cta 8270  
 Met Glu Lys Ile Glu Lys Phe Lys Asn Lys Cys Gln His Lys Leu  
 1 5 10 15  
 caa cat aaa cta atc gta tta gta tca aca ctt tgc tat ata aac aat 8318  
 Gln His Lys Leu Ile Val Leu Val Ser Thr Leu Cys Tyr Ile Asn Asn  
 20 25 30  
 aaa aat aaa aaa tat tca caa agc aac atc ctt tat tat ttt aat gaa 8366  
 Lys Asn Lys Lys Tyr Ser Gln Ser Asn Ile Leu Tyr Tyr Phe Asn Glu  
 35 40 45  
 aat tta aaa aga aat ggg caa acc cct att aaa ata aaa aca tta caa 8414  
 Asn Leu Lys Arg Asn Gly Gln Thr Pro Ile Lys Ile Lys Thr Leu Gln  
 50 55 60  
 aac tat ctt tat aaa ctg gaa aaa gaa ttt gaa gta aca act aat tat 8462  
 Asn Tyr Leu Tyr Lys Leu Glu Lys Glu Phe Glu Val Thr Thr Asn Tyr  
 65 70 75  
 tat aaa cac ttg ggg gtt aat tgt gga acc gaa att tac tat aaa ctt 8510  
 Tyr Lys His Leu Gly Val Asn Cys Gly Thr Glu Ile Tyr Tyr Lys Leu  
 80 85 90 95  
 aaa tat caa aaa caa aaa tgc tat cat aaa ata aac caa tat ttt aaa 8558  
 Lys Tyr Gln Lys Gln Lys Cys Tyr His Lys Ile Asn Gln Tyr Phe Lys  
 100 105 110  
 aag aaa aaa gaa att aaa ttt aac tta aga gta agt gca ttt ttt aat 8606  
 Lys Lys Lys Glu Ile Lys Phe Asn Leu Arg Val Ser Ala Phe Phe Asn  
 115 120 125  
 aaa aaa cac tca aaa aaa ggg agt gta gaa tta aag gaa tgt aat aat 8654  
 Lys Lys His Ser Lys Lys Gly Ser Val Glu Leu Lys Glu Cys Asn Asn  
 130 135 140  
 aat aat aat aat aaa gag aaa gaa aca tcc caa aaa att gaa att tta 8702

Asn Asn Asn Asn Lys Glu Lys Glu Thr Ser Gln Lys Ile Glu Ile Leu  
 145 150 155

caa aca aaa gtc tat gcc aaa aaa tgt aaa ttt ttg aca aac tac tat 8750  
 Gln Thr Lys Val Tyr Ala Lys Lys Cys Lys Phe Leu Thr Asn Tyr Tyr  
 160 165 170 175

act aaa att tta 8762  
 Thr Lys Ile Leu

<210> 6  
 <211> 179  
 <212> PRT  
 <213> Borrelia afzelii

<400> 6  
 Met Glu Lys Ile Glu Lys Phe Lys Asn Lys Cys Gln His Lys Leu Gln  
 1 5 10 15

His Lys Leu Ile Val Leu Val Ser Thr Leu Cys Tyr Ile Asn Asn Lys  
 20 25 30

Asn Lys Lys Tyr Ser Gln Ser Asn Ile Leu Tyr Tyr Phe Asn Glu Asn  
 35 40 45

Leu Lys Arg Asn Gly Gln Thr Pro Ile Lys Ile Lys Thr Leu Gln Asn  
 50 55 60

Tyr Leu Tyr Lys Leu Glu Lys Glu Phe Glu Val Thr Thr Asn Tyr Tyr  
 65 70 75 80

Lys His Leu Gly Val Asn Cys Gly Thr Glu Ile Tyr Tyr Lys Leu Lys  
 85 90 95

Tyr Gln Lys Gln Lys Cys Tyr His Lys Ile Asn Gln Tyr Phe Lys Lys  
 100 105 110

Lys Lys Glu Ile Lys Phe Asn Leu Arg Val Ser Ala Phe Phe Asn Lys  
 115 120 125

Lys His Ser Lys Lys Gly Ser Val Glu Leu Lys Glu Cys Asn Asn Asn  
 130 135 140

Asn Asn Asn Lys Glu Lys Glu Thr Ser Gln Lys Ile Glu Ile Leu Gln  
 145 150 155 160

Thr Lys Val Tyr Ala Lys Lys Cys Lys Phe Leu Thr Asn Tyr Tyr Thr  
 165 170 175

Lys Ile Leu

<210> 7  
 <211> 416

<212> DNA

<213> *Borrelia afzelii*

<220>

<221> CDS

<222> (1)..(414)

<400> 7

aag	ggg	att	gcg	aag	ggg	ata	aag	ggg	att	gtt	gcg	gct	gct	ggg	aag	48
Lys	Gly	Ile	Ala	Lys	Gly	Ile	Lys	Gly	Ile	Val	Ala	Ala	Ala	Gly	Lys	
1				5				10						15		

gct	ttt	ggc	aag	gat	ggg	gat	gcg	ctg	aca	ggg	gtt	gca	aaa	gct	gct	96
Ala	Phe	Gly	Lys	Asp	Gly	Asp	Ala	Leu	Thr	Gly	Val	Ala	Lys	Ala	Ala	
		20					25					30				

gag	aat	gat	gct	aac	aag	gat	gcg	ggg	aag	ttg	ttt	gct	ggg	aag	aat	144
Glu	Asn	Asp	Ala	Asn	Lys	Asp	Ala	Gly	Lys	Leu	Phe	Ala	Gly	Lys	Asn	
		35					40					45				

ggg	aat	gct	ggg	gct	gct	gac	att	gcg	aag	gcg	gct	gct	gct	gct	gtt	act	192
Gly	Asn	Ala	Gly	Ala	Ala	Asp	Ile	Ala	Lys	Ala	Ala	Ala	Ala	Ala	Val	Thr	
	50					55				60							

gcg	gtt	agt	ggg	gag	cag	ata	cta	aaa	gct	att	gtt	gag	gcg	gct	ggg	240
Ala	Val	Ser	Gly	Glu	Gln	Ile	Leu	Lys	Ala	Ile	Val	Glu	Ala	Ala	Gly	
65				70				75							80	

gat	gcg	gat	cag	gcg	ggg	gta	aag	gct	gat	gcg	gct	aag	aat	ccg	att	288
Asp	Ala	Asp	Gln	Ala	Gly	Val	Lys	Ala	Asp	Ala	Ala	Lys	Asn	Pro	Ile	
			85					90						95		

gca	gct	gcg	att	ggg	act	gct	gat	gat	ggg	gct	gcg	ttt	ggg	aag	gat	336
Ala	Ala	Ala	Ile	Gly	Thr	Ala	Asp	Asp	Gly	Ala	Ala	Phe	Gly	Lys	Asp	
			100					105					110			

gag	atg	aag	aag	aga	aat	gat	aag	att	gtt	gca	gct	att	gtt	ttg	agg	384
Glu	Met	Lys	Lys	Arg	Asn	Asp	Lys	Ile	Val	Ala	Ala	Ile	Val	Leu	Arg	
		115					120					125				

ggg	gtg	cct	aag	gat	gga	aag	ttt	gct	gct	aa						416
Gly	Val	Pro	Lys	Asp	Gly	Lys	Phe	Ala	Ala							
	130					135										

<210> 8

<211> 138

<212> PRT

<213> *Borrelia afzelii*

<400> 8

Lys	Gly	Ile	Ala	Lys	Gly	Ile	Lys	Gly	Ile	Val	Ala	Ala	Ala	Gly	Lys	
1				5				10						15		

Ala	Phe	Gly	Lys	Asp	Gly	Asp	Ala	Leu	Thr	Gly	Val	Ala	Lys	Ala	Ala	
		20					25						30			



att gcg gct gcg att ggg act gct gat gat ggg gcg gag ttt aag gat 336  
 Ile Ala Ala Ala Ile Gly Thr Ala Asp Asp Gly Ala Glu Phe Lys Asp  
                   100                  105                  110

gat atg aag aag agt gat aat att gct gcg gct att gtt ttg agg ggg 384  
 Asp Met Lys Lys Ser Asp Asn Ile Ala Ala Ala Ile Val Leu Arg Gly  
                   115                  120                  125

gtg cct aag gat gga aag ttt gct gct aa 413  
 Val Pro Lys Asp Gly Lys Phe Ala Ala  
                   130                  135

<210> 10  
 <211> 137  
 <212> PRT  
 <213> Borrelia afzelii

<400> 10  
 Lys Gly Ile Ala Lys Gly Ile Lys Gly Ile Val Asp Ala Ala Gly Lys  
   1                  5                  10                  15  
 Ala Phe Gly Lys Glu Gly Ser Ala Leu Lys Asp Val Ala Lys Val Ala  
                   20                  25                  30  
 Asp Asp Asp Asn Lys Asp Ala Gly Lys Leu Phe Ala Gly Lys Asn Gly  
                   35                  40                  45  
 Gly Ala Gly Ala Ala Asp Ala Ile Gly Lys Ala Ala Ala Val Thr  
   50                  55                  60  
 Ala Val Ser Gly Glu Gln Ile Leu Lys Ala Ile Val Asp Ala Ala Gly  
   65                  70                  75                  80  
 Ala Ala Ala Asn Gln Ala Gly Lys Lys Ala Ala Asp Ala Lys Asn Pro  
                   85                  90                  95  
 Ile Ala Ala Ala Ile Gly Thr Ala Asp Asp Gly Ala Glu Phe Lys Asp  
                   100                  105                  110  
 Asp Met Lys Lys Ser Asp Asn Ile Ala Ala Ala Ile Val Leu Arg Gly  
                   115                  120                  125  
 Val Pro Lys Asp Gly Lys Phe Ala Ala  
   130                  135

<210> 11  
 <211> 428  
 <212> DNA  
 <213> Borrelia afzelii

<220>  
 <221> CDS  
 <222> (1)..(426)



<400> 11

aag ggg att gcg aag ggg ata aag ggg att gtt gat gct gct ggg aag 48  
Lys Gly Ile Ala Lys Gly Ile Lys Gly Ile Val Asp Ala Ala Gly Lys  
1 5 10 15

gct ttt ggt aag gag ggt gat gcg ctg aag gat gtt gca aaa gtt gct 96  
Ala Phe Gly Lys Glu Gly Asp Ala Leu Lys Asp Val Ala Lys Val Ala  
20 25 30

gat gag aat ggg gat aac aag gat gcg ggg aag ttg ttt gct ggt gag 144  
Asp Glu Asn Gly Asp Asn Lys Asp Ala Gly Lys Leu Phe Ala Gly Glu  
35 40 45

aat ggt aat gct ggt ggt gct gct gat gct gac att gcg aag gcg gct 192  
Asn Gly Asn Ala Gly Gly Ala Ala Asp Ala Asp Ile Ala Lys Ala Ala  
50 55 60

gct gct gtt act gcg gtt agt ggg gag cag ata ctg aaa gct att gtt 240  
Ala Ala Val Thr Ala Val Ser Gly Glu Gln Ile Leu Lys Ala Ile Val  
65 70 75 80

gag gcg gct ggt gct gcg gat cag gcg ggt gta aag gct gag gag gct 288  
Glu Ala Ala Gly Ala Ala Asp Gln Ala Gly Val Lys Ala Glu Glu Ala  
85 90 95

aag aat ccg att gca gct gcg att ggg act gat gat gct ggt gcg gcg 336  
Lys Asn Pro Ile Ala Ala Ala Ile Gly Thr Asp Asp Ala Gly Ala Ala  
100 105 110

gag ttt ggt gag aat gat atg aag aag aat gat aat att gct gcg gct 384  
Glu Phe Gly Glu Asn Asp Met Lys Lys Asn Asp Asn Ile Ala Ala Ala  
115 120 125

att gtt ttg agg ggg gtg cct aag gat gga aag ttt gct gct aa 428  
Ile Val Leu Arg Gly Val Pro Lys Asp Gly Lys Phe Ala Ala  
130 135 140

<210> 12

<211> 142

<212> PRT

<213> *Borrelia afzelii*

<400> 12

Lys Gly Ile Ala Lys Gly Ile Lys Gly Ile Val Asp Ala Ala Gly Lys  
1 5 10 15

Ala Phe Gly Lys Glu Gly Asp Ala Leu Lys Asp Val Ala Lys Val Ala  
20 25 30

Asp Glu Asn Gly Asp Asn Lys Asp Ala Gly Lys Leu Phe Ala Gly Glu  
35 40 45

Asn Gly Asn Ala Gly Gly Ala Ala Asp Ala Asp Ile Ala Lys Ala Ala  
50 55 60

Ala Ala Val Thr Ala Val Ser Gly Glu Gln Ile Leu Lys Ala Ile Val

65		70		75		80									
Glu	Ala	Ala	Gly	Ala	Ala	Asp	Gln	Ala	Gly	Val	Lys	Ala	Glu	Glu	Ala
			85						90					95	
Lys	Asn	Pro	Ile	Ala	Ala	Ala	Ile	Gly	Thr	Asp	Asp	Ala	Gly	Ala	Ala
			100					105					110		
Glu	Phe	Gly	Glu	Asn	Asp	Met	Lys	Lys	Asn	Asp	Asn	Ile	Ala	Ala	Ala
		115					120					125			
Ile	Val	Leu	Arg	Gly	Val	Pro	Lys	Asp	Gly	Lys	Phe	Ala	Ala		
	130					135					140				

<210> 13  
 <211> 426  
 <212> DNA  
 <213> *Borrelia afzelii*

<220>  
 <221> CDS  
 <222> (3) .. (425)

<400> 13	
ag ggg att gcg aag ggg ata aag ggg att gtt gat gct gct ggg aag	47
Gly Ile Ala Lys Gly Ile Lys Gly Ile Val Asp Ala Ala Gly Lys	
1 5 10 15	
gct ttt ggc aag gag ggt agt gcg ctg aag gat gtt aaa aca gtt gct	95
Ala Phe Gly Lys Glu Gly Ser Ala Leu Lys Asp Val Lys Thr Val Ala	
20 25 30	
gct gag aat gag gct aac aag gat gcg ggg aag ttg ttt gct ggt aag	143
Ala Glu Asn Glu Ala Asn Lys Asp Ala Gly Lys Leu Phe Ala Gly Lys	
35 40 45	
aat ggt aat gct gat gct gct gat gct gct gac att gcg aag gcg gct	191
Asn Gly Asn Ala Asp Ala Ala Asp Ala Ala Asp Ile Ala Lys Ala Ala	
50 55 60	
ggt gct gtt agt gcg gtt agt ggg gag cag ata ctg aaa gct att gtt	239
Gly Ala Val Ser Ala Val Ser Gly Glu Gln Ile Leu Lys Ala Ile Val	
65 70 75	
gat ggt gct ggt gat gca gct aat cag gcg ggt aaa aag gct gct gag	287
Asp Gly Ala Gly Asp Ala Ala Asn Gln Ala Gly Lys Lys Ala Ala Glu	
80 85 90 95	
gct aag aat ccg att gcg gct gcg att ggg act aat gaa gct ggg gcg	335
Ala Lys Asn Pro Ile Ala Ala Ala Ile Gly Thr Asn Glu Ala Gly Ala	
100 105 110	
gag ttt ggt gat gat atg aag aag aga aat gat aag att gct gcg gct	383
Glu Phe Gly Asp Asp Met Lys Lys Arg Asn Asp Lys Ile Ala Ala Ala	
115 120 125	

att gtt ttg agg ggg gtg cct aag gat gga aag ttt gct gct a 426  
 Ile Val Leu Arg Gly Val Pro Lys Asp Gly Lys Phe Ala Ala  
           130                          135                          140

<210> 14  
 <211> 141  
 <212> PRT  
 <213> *Borrelia afzelii*

<400> 14  
 Gly Ile Ala Lys Gly Ile Lys Gly Ile Val Asp Ala Ala Gly Lys Ala  
       1                          5                          10                          15  
 Phe Gly Lys Glu Gly Ser Ala Leu Lys Asp Val Lys Thr Val Ala Ala  
                           20                          25                          30  
 Glu Asn Glu Ala Asn Lys Asp Ala Gly Lys Leu Phe Ala Gly Lys Asn  
                           35                          40                          45  
 Gly Asn Ala Asp Ala Ala Asp Ala Ala Asp Ile Ala Lys Ala Ala Gly  
                           50                          55                          60  
 Ala Val Ser Ala Val Ser Gly Glu Gln Ile Leu Lys Ala Ile Val Asp  
                           65                          70                          75                          80  
 Gly Ala Gly Asp Ala Ala Asn Gln Ala Gly Lys Lys Ala Ala Glu Ala  
                           85                          90                          95  
 Lys Asn Pro Ile Ala Ala Ala Ile Gly Thr Asn Glu Ala Gly Ala Glu  
                           100                          105                          110  
 Phe Gly Asp Asp Met Lys Lys Arg Asn Asp Lys Ile Ala Ala Ala Ile  
                           115                          120                          125  
 Val Leu Arg Gly Val Pro Lys Asp Gly Lys Phe Ala Ala  
                           130                          135                          140

<210> 15  
 <211> 5897  
 <212> DNA  
 <213> *Borrelia garinii*

<400> 15  
 cggaaatcaa gccacctaaa acaacttccc aaaagtttct caaaaaatat tatattcagc 60  
 agtaaattct ataagtcatt aattatttaa tactattcaa cagtaaattc tataagtcac 120  
 taattattta atactattca gcagtaaatt ctataagtca ttaattattt aatactattc 180  
 agcagtaaatt tctataagtc attaattatt taatactatt cagcagtaaa ttctataagt 240  
 cattaattat ttaatactat tcagcagtaa attctataag tcattaatta ttttaacta 300  
 ttcagcagta aattctataa gtcattaatt caattaggtta acggattctt agatgtattc 360  
 acctcttttg gtggattagt tgcagatgca ttgggggtta aagctgatcc aaaaaaatct 420  
 gatgtaaaaa cttattttga atctctagct aaaaaattag aagaaacaaa agatgggttta 480  
 actaagttgt ccaaaggtaa tgacggtgat actggaaagg ctggtgatgc tgggtggggct 540  
 ggtggtggcg ctagtgctgc aggtggcgct ggtgggattg agggcgctat aacagagatt 600

agcaaatggt	tagatgatat	ggcaaaagct	gctgcggaag	ctgcaagtgc	tgctactggt	660
aatgcagcaa	ttggggatgt	tgtaaatggt	aatggtggag	cagcaaaagg	tggtgatgctg	720
gagagtgtta	atgggattgc	taaggggata	aaggggattg	ttgatgctgc	tgagaaggct	780
gatgcgaagg	aagggaagtt	ggatgtggct	ggtgatgctg	gtggggctgg	tggtggcgct	840
ggtgctgcag	gtggcgctgg	tgggattgag	ggcgctataa	cagagattag	caaatggtta	900
gatgatatgg	caaaagctgc	tgcggttgct	gcaagtgctg	caagtgctgc	tactggtaat	960
gcagcaattg	gggatgttgt	taatggtaat	gatggagcag	caaaagggtg	tgatgcggcg	1020
agtgttaatg	ggattgctaa	ggggataaa	gggattgttg	atgctgctga	gaaggctgat	1080
gcgaaggaa	ggaagtggga	tgtggctggt	gatgctggtg	agggtacaa	ggatgctggg	1140
aagctgtttg	tgaagaagaa	tgctggtgat	gaggggtggtg	aagcaaatga	tgctgggaag	1200
gctgctgctg	cggttgctgc	tgtagtggg	gagcagatat	taaaagcgat	tgtagtgc	1260
gctgaggggtg	atgataagca	gggtaagaag	gctgcggatg	ctacaaatcc	gattgaggcg	1320
gctattgggg	gtgcggatgc	gggtgcta	gctgaggcgt	ttaataagat	gaagaaggat	1380
gatcagattg	ctgctgctat	ggttctgagg	ggaatggcta	aggatgggca	gtttgctttg	1440
aaggatgatg	ctgctgctca	tgaagggact	gttaagaatg	ctgttgatat	ggcaaaggcc	1500
gctgcggaag	ctgcaagtgc	tgcaagtgct	gctactggta	gtacaacgat	tgagatggt	1560
gttaagagtg	gtgaggcaaa	agatggtgat	gcggcgagtg	ttaatgggat	tgctaagggg	1620
ataaagggga	ttgttgatgc	tgctgagaag	gctgatgcga	aggaaggga	gttgatgtg	1680
gctggtgctg	ctggtacgac	taacgtgaat	gttgggaagt	tgtagtgaa	gaataatggt	1740
aatgaggggtg	gtgatgcaag	tgatgctggg	aaagctgctg	ctgcggttgc	tgctgttagt	1800
ggggagcaga	tattaaaagc	gattgttgat	gctgctaagg	atggtgataa	gcagggggtt	1860
actgatgtaa	aggatgctac	aatccgatt	gaggcggcta	ttgggggtac	aatgataat	1920
gatgctgcgg	cgtttgctac	tatgaagaag	gatgatcaga	ttgctgctgc	tatggttctg	1980
aggggaatgg	ctaaggatgg	gcagtttgct	ttgaaggatg	atgctgctaa	ggatggtgat	2040
aaaacggggg	ttgctgcgga	tgctgaaaat	ccgattgacg	cggctattgg	gggtgcggat	2100
gctgatgctg	cggcgtttaa	taaggagggg	atgaagaagg	atgatcagat	tgctgctgct	2160
gtggttctga	ggggaatggc	taaggatggg	cagtttgctt	tgacgaataa	tgctgctgct	2220
catgaaggga	ctgttaagaa	tgctgttgat	atggcaaaag	ctgctgcggt	tgctgcaagt	2280
gctgctactg	gcaatgcagc	aattggggat	gttgttaaga	gtaatggtgg	agcagcagca	2340
aaaggtgggtg	atgcggcgag	tgtaaatggg	attgctaagg	ggataaagg	gattgttgat	2400
gctgctgaga	aggctgatgc	gaagggaagg	aagtggatg	tggtggtgc	tgctggtgaa	2460
actaacaagg	atgctgggaa	gttgtttgtg	aagaagaatg	gtgatgatgg	tggtgatgca	2520
ggtgatgctg	ggaaggctgc	tgctgcggtt	gctgctgtta	gtggggagca	gatattaaaa	2580
gcgattgttg	atgctgctaa	agatggtgat	aagacggggg	ttactgatgt	aaaggatgct	2640
acaaatccga	ttgacgcggc	tattgggggg	agtgcggatg	ctaagtctga	ggcgtttgat	2700
aagatgaaga	aggatgatca	gattgctgct	gctatggttc	tgaggggaat	ggctaaggat	2760
gggcagtttg	ctttgaagaa	taatgatcat	gataatcata	aggggactgt	taagaatgct	2820
gttgatatgg	caaaggccgc	tgaggaagct	gcaagtgctg	caagtgctgc	tactggtaat	2880
gcagcgattg	gggatgttgt	taagaatagt	ggggcagcag	caaaagggtg	tgaggcggcg	2940
agtgttaatg	ggattgctaa	ggggataaa	gggattgttg	atgctgctgg	aaaggctgat	3000
gcgaaggaa	ggaagtggga	tgctactggt	gctgagggtg	cgactaacgt	gaatgctggg	3060
aagttgtttg	tgaagagggc	ggctgatgat	ggtggtgatg	cagatgatgc	tggaaggct	3120
gctgctgcgg	ttgctgcaag	tgctgctact	ggtaatgcag	cgattggaga	tgtagttaa	3180
ggtgatgtgg	caaaagcaaa	aggtggtgat	gcggcgagtg	ttaatgggat	tgctaagggg	3240
ataaagggga	ttgttgatgc	tgctgagaag	gctgatgcga	aggaaggga	gttgatgct	3300
gctggtgctg	aggggtacgac	taacgcggat	gctgggaagt	tgtagtgaa	gaatgctggt	3360
aatgtgggtg	gtgaagcagg	tgatgctggg	aaggctgctg	ctgcggttgc	tgctgttagt	3420
ggggagcaga	tattaaaagc	gattgttgat	gctgctaagg	atggtggtga	gaagcagggt	3480
aagaaggctg	cggatgctac	aatccgatt	gacgcggcta	ttgggggtac	aatgataat	3540
gatgctgctg	cggcgtttgc	tactatgaag	aaggatgatc	agattgctgc	tgctatggtt	3600
ctgaggggaa	tggtctaa	tggtcaattt	gctttgaagg	atgctgctgc	tgctcatgaa	3660
gggactgtta	agaatgctgt	tgatataata	aaggctgctg	cggaagctgc	aagtgctgca	3720
agtgctgcta	ctggtagtgc	agcaattggg	gatgttggtta	atggtaatgg	agcaacagca	3780
aaaggtgggtg	atgcgaagag	tgtaaatggg	attgctaagg	ggataaagg	gattgttgat	3840
gctgctgaga	aggctgatgc	gaagggaagg	aagtggatg	tggtggtga	tgctggtgaa	3900
actaacaagg	atgctgggaa	gttgtttgtg	aagaacaatg	gtaatgagg	tggtgatgca	3960
gatgatgctg	ggaaggctgc	tgctgcggtt	gctgctgtta	gtggggagca	gatattaaaa	4020

```

gcgattgttg atgctgctaa ggggtggtgat aagacgggta agaataatgt gaaggatgct 4080
gaaaatccga ttgaggcggc tattgggagt agtgcggatg ctgatgctgc ggcgtttaat 4140
aaggagggga tgaagaagga tgatcagatt gctgctgcta tggttctgag gggaatggct 4200
aaggatgggc agtttgcttt gacgaatgat gctgctgctc atgaagggac tgtaagaat 4260
gctgttggga gtgcaacaaa taagaccgtt gttgcttttg ctaacttggg tcgaaagacc 4320
gtgcaagctg ggttgaagaa ggttggggat gttgttaaga atagtgaggc aaaagatggt 4380
gatgcggcga gtgttaatgg gattgctaag gggataaagg ggattgttga tgctgctgag 4440
aaggctgatg cgaaggaagg gaagttggat gtggctgggt ctgctgggtga aactaacaag 4500
gatgctggga agttgtttgt gaagaagaat aatgaggggt gtgaagcaaa tgatgctggg 4560
aaggctgctg ctgcggttgc tgctgttagt ggggagcaga tattaaaagc gattgttgat 4620
gctgctaagg atggtgatga taagcagggt aagaaggctg aggatgctac aaatccgatt 4680
gacgcggcta ttgggggtgc aggtgcgggt gctaattgctg ctgcggcggt taataatatg 4740
aagaaggatg atcagattgc tgctgctatg gttctgaggg gaatggctaa ggatgggcag 4800
tttgctttga cgaataatgc tcataactaat cataagggga ctgttaagaa tgctgttgat 4860
atgacaaaag ctgctgcggt tgctgcaagt gctgcaagtg ctgctactgg taatgcagca 4920
attggggatg ttgttaatgg taatgatgga gcagcaaaaag gtggtgatgc ggcgagtgtt 4980
aatgggattg ctaaggggat aaaggggatt gttgatgctg ctgagaaggc tgatgcgaag 5040
gaagggaagt tgaatgtggc tgggtgctgct ggtgctgagg gtaacgaggc tgctgggaag 5100
ctgtttgtga agaagaatgc tggtgatcat ggtggtgaag caggtgatgc tgggagggct 5160
gctgctgcgg ttgctgctgt tagtggggag cagatattaa aagcgattgt tgatgctgct 5220
aaggatggtg gtgataagca gggtaagaag gctgaggatg ctgaaaatcc gattgacgcg 5280
gctattggga gtacgggtgc ggatgataat gctgctgagg cgtttgctac tatgaagaag 5340
gatgatcaga ttgctgctgc tatggttctg aggggaatgg ctaaggatgg gcagtttgct 5400
ttgaaggatg ctgctcatga taatcataag gggactgtta agaatgctgt tgatataata 5460
aaggctactg cggttgctgc aagtgctgct actggtagta caacgattgg ggatgttggt 5520
aagaatggtg aggcaaaaagg tggtagggcg aagagtgtta atgggattgc taaggggata 5580
aaggggattg ttgatgctgc tggaaaaggct gatgcgaagg aagggaaagt gaatgtggct 5640
ggtgctgctg tgtacggtaa cgaggctgct gggaaagctgt ttgtgtaaat tactatagga 5700
ttagaactag gtgacgatat gagtcccttg gttattttgc agctgctaata gaatttgaaa 5760
taagtgaagt taaaattgcg gatgttaatg gaacacattt tattgctaca aaagagaaaag 5820
aaatattata tgattcactt gatttaagggt ctctgaggaa aatatttgaa ataacttcaa 5880
agcgaatggt taagctt 5897

```

<210> 16  
 <211> 396  
 <212> DNA  
 <213> *Borrelia garinii*

<220>  
 <221> CDS  
 <222> (2) .. (394)

```

<400> 16
g ggg ata aag ggg att gtt gat gct gct gag aag gct gat gcg aag gaa 49
  Gly Ile Lys Gly Ile Val Asp Ala Ala Glu Lys Ala Asp Ala Lys Glu
    1             5             10             15

ggg aag ttg aat gct gct ggt gct gag ggt acg act aac gcg gat gct 97
Gly Lys Leu Asn Ala Ala Gly Ala Glu Gly Thr Thr Asn Ala Asp Ala
    20             25             30

ggg aag ttg ttt gtg aag aat gct ggt aat gtg ggt ggt gaa gca ggt 145
Gly Lys Leu Phe Val Lys Asn Ala Gly Asn Val Gly Gly Glu Ala Gly
    35             40             45

gat gct ggg aag gct gct gct gcg gtt gct gct gtt agt ggg gag cag 193

```



<210> 18  
 <211> 390  
 <212> DNA  
 <213> *Borrelia garinii*

<220>  
 <221> CDS  
 <222> (2) .. (388)

<400> 18  
 g ggg ata aag ggg att gtt gat gct gct gag aag gct gat gcg aag gaa 49  
 Gly Ile Lys Gly Ile Val Asp Ala Ala Glu Lys Ala Asp Ala Lys Glu  
 1 5 10 15

ggg aag ttg gat gtg gct ggt gat gct ggt gaa act aac aag gat gct 97  
 Gly Lys Leu Asp Val Ala Gly Asp Ala Gly Glu Thr Asn Lys Asp Ala  
 20 25 30

ggg aag ttg ttt gtg aag aag aat aat gag ggt ggt gaa gca aat gat 145  
 Gly Lys Leu Phe Val Lys Lys Asn Asn Glu Gly Gly Glu Ala Asn Asp  
 35 40 45

gct ggg aag gct gct gct gcg gtt gct gct gtt agt ggg gag cag ata 193  
 Ala Gly Lys Ala Ala Ala Val Ala Ala Val Ser Gly Glu Gln Ile  
 50 55 60

tta aaa gcg att gtt gat gct gct gag ggt ggt gag aag cag ggt aag 241  
 Leu Lys Ala Ile Val Asp Ala Ala Glu Gly Gly Glu Lys Gln Gly Lys  
 65 70 75 80

aag gct gcg gat gct aca aat ccg att gag gcg gct att ggg ggt gcg 289  
 Lys Ala Ala Asp Ala Thr Asn Pro Ile Glu Ala Ala Ile Gly Gly Ala  
 85 90 95

ggt gat aat gat gct gct gcg gcg ttt gct act atg aag aag gat gat 337  
 Gly Asp Asn Asp Ala Ala Ala Ala Phe Ala Thr Met Lys Lys Asp Asp  
 100 105 110

cag att gct act gct atg gtt ctg agg gga atg gct aag gat ggg cag 385  
 Gln Ile Ala Thr Ala Met Val Leu Arg Gly Met Ala Lys Asp Gly Gln  
 115 120 125

ttt gc 390  
 Phe

<210> 19  
 <211> 129  
 <212> PRT  
 <213> *Borrelia garinii*

<400> 19  
 Gly Ile Lys Gly Ile Val Asp Ala Ala Glu Lys Ala Asp Ala Lys Glu  
 1 5 10 15

Gly Lys Leu Asp Val Ala Gly Asp Ala Gly Glu Thr Asn Lys Asp Ala  
                   20                                  25                                  30  
 Gly Lys Leu Phe Val Lys Lys Asn Asn Glu Gly Gly Glu Ala Asn Asp  
                   35                                  40                                  45  
 Ala Gly Lys Ala Ala Ala Ala Val Ala Ala Val Ser Gly Glu Gln Ile  
                   50                                  55                                  60  
 Leu Lys Ala Ile Val Asp Ala Ala Glu Gly Gly Glu Lys Gln Gly Lys  
                   65                                  70                                  75                                  80  
 Lys Ala Ala Asp Ala Thr Asn Pro Ile Glu Ala Ala Ile Gly Gly Ala  
                                   85                                  90                                  95  
 Gly Asp Asn Asp Ala Ala Ala Ala Phe Ala Thr Met Lys Lys Asp Asp  
                                   100                                  105                                  110  
 Gln Ile Ala Thr Ala Met Val Leu Arg Gly Met Ala Lys Asp Gly Gln  
                   115                                  120                                  125

Phe

<210> 20  
 <211> 390  
 <212> DNA  
 <213> *Borrelia garinii*

<220>  
 <221> CDS  
 <222> (2) .. (388)

<400> 20  
 g ggg ata aag ggg att gtt gat gct gct gag aag gct gat gcg aag gaa 49  
   Gly Ile Lys Gly Ile Val Asp Ala Ala Glu Lys Ala Asp Ala Lys Glu  
       1                                  5                                  10                                  15  
  
 ggg agg ttg gat gtg gct ggt gat gct ggt gaa act aac aag gat gct 97  
   Gly Arg Leu Asp Val Ala Gly Asp Ala Gly Glu Thr Asn Lys Asp Ala  
                   20                                  25                                  30  
  
 ggg aag ttg ttt gtg aag aag aat aat gag ggt ggt gaa gca aat gat 145  
   Gly Lys Leu Phe Val Lys Lys Asn Asn Glu Gly Gly Glu Ala Asn Asp  
                   35                                  40                                  45  
  
 gct ggg aag gct gct gct gcg gtt gct gct gtt agt ggg gag cag ata 193  
   Ala Gly Lys Ala Ala Ala Ala Val Ala Ala Val Ser Gly Glu Gln Ile  
                   50                                  55                                  60  
  
 tta aaa gcg att gtt gat gct gct gag ggt ggt gag aag cag ggt aag 241  
   Leu Lys Ala Ile Val Asp Ala Ala Glu Gly Gly Glu Lys Gln Gly Lys  
                   65                                  70                                  75                                  80  
  
 aag gct gcg gat gct aca aat ccg att gag gcg gct att ggg ggt gcg 289



Lys	Ala	Ala	Asp	Ala	Thr	Asn	Pro	Ile	Glu	Ala	Ala	Ile	Gly	Gly	Ala	
				85					90					95		
ggt	gat	aat	gat	gct	gct	gcg	gcg	ttt	gct	act	atg	aag	aag	gat	gat	337
Gly	Asp	Asn	Asp	Ala	Ala	Ala	Ala	Phe	Ala	Thr	Met	Lys	Lys	Asp	Asp	
			100					105					110			
cag	att	gct	gct	gct	atg	gtt	ctg	agg	gga	atg	gct	aag	gat	ggg	cag	385
Gln	Ile	Ala	Ala	Ala	Met	Val	Leu	Arg	Gly	Met	Ala	Lys	Asp	Gly	Gln	
			115				120					125				
ttt	gc															390
Phe																

<210> 21  
 <211> 129  
 <212> PRT  
 <213> *Borrelia garinii*

Gly	Ile	Lys	Gly	Ile	Val	Asp	Ala	Ala	Glu	Lys	Ala	Asp	Ala	Lys	Glu	
1				5					10					15		
Gly	Arg	Leu	Asp	Val	Ala	Gly	Asp	Ala	Gly	Glu	Thr	Asn	Lys	Asp	Ala	
			20					25					30			
Gly	Lys	Leu	Phe	Val	Lys	Lys	Asn	Asn	Glu	Gly	Gly	Glu	Ala	Asn	Asp	
		35					40					45				
Ala	Gly	Lys	Ala	Ala	Ala	Ala	Val	Ala	Ala	Val	Ser	Gly	Glu	Gln	Ile	
	50					55					60					
Leu	Lys	Ala	Ile	Val	Asp	Ala	Ala	Glu	Gly	Gly	Glu	Lys	Gln	Gly	Lys	
65					70				75					80		
Lys	Ala	Ala	Asp	Ala	Thr	Asn	Pro	Ile	Glu	Ala	Ala	Ile	Gly	Gly	Ala	
				85					90					95		
Gly	Asp	Asn	Asp	Ala	Ala	Ala	Ala	Phe	Ala	Thr	Met	Lys	Lys	Asp	Asp	
			100					105					110			
Gln	Ile	Ala	Ala	Ala	Met	Val	Leu	Arg	Gly	Met	Ala	Lys	Asp	Gly	Gln	
		115					120					125				
Phe																

<210> 22  
 <211> 339  
 <212> DNA  
 <213> *Borrelia garinii*

<220>  
 <221> CDS

<222> (2) .. (337)

<400> 22

```
g ggg ata aag ggg att gtt gat gct gct ggt gaa act aac aag gat gct 49
  Gly Ile Lys Gly Ile Val Asp Ala Ala Gly Glu Thr Asn Lys Asp Ala
    1             5             10             15

ggg aag ttg ttt gtg aag aag aat aat gag ggt ggt gaa gca aat gat 97
  Gly Lys Leu Phe Val Lys Lys Asn Asn Glu Gly Gly Glu Ala Asn Asp
    20             25             30

gct ggg aag gct gct gct gcg gtt gct gct gtt agt ggg gag cag ata 145
  Ala Gly Lys Ala Ala Ala Ala Val Ala Ala Val Ser Gly Glu Gln Ile
    35             40             45

tta aaa gcg att gtt gat gct gct gag ggt ggt gag aag cag ggt aag 193
  Leu Lys Ala Ile Val Asp Ala Ala Glu Gly Gly Glu Lys Gln Gly Lys
    50             55             60

aag gct gcg gat gct aca aat ccg att gag gcg gct att ggg ggt aca 241
  Lys Ala Ala Asp Ala Thr Asn Pro Ile Glu Ala Ala Ile Gly Gly Thr
    65             70             75             80

aat gat aat gat gct gcg gcg ttt gct act atg aag aag gat gat cag 289
  Asn Asp Asn Asp Ala Ala Ala Phe Ala Thr Met Lys Lys Asp Asp Gln
    85             90             95

att gct gct gct atg gtt ctg agg gga atg gct aag gat ggg cag ttt 337
  Ile Ala Ala Ala Met Val Leu Arg Gly Met Ala Lys Asp Gly Gln Phe
    100            105            110

gc 339
```

<210> 23

<211> 112

<212> PRT

<213> *Borrelia garinii*

<400> 23

```
Gly Ile Lys Gly Ile Val Asp Ala Ala Gly Glu Thr Asn Lys Asp Ala
  1             5             10             15

Gly Lys Leu Phe Val Lys Lys Asn Asn Glu Gly Gly Glu Ala Asn Asp
  20             25             30

Ala Gly Lys Ala Ala Ala Ala Val Ala Ala Val Ser Gly Glu Gln Ile
  35             40             45

Leu Lys Ala Ile Val Asp Ala Ala Glu Gly Gly Glu Lys Gln Gly Lys
  50             55             60

Lys Ala Ala Asp Ala Thr Asn Pro Ile Glu Ala Ala Ile Gly Gly Thr
  65             70             75             80

Asn Asp Asn Asp Ala Ala Ala Phe Ala Thr Met Lys Lys Asp Asp Gln
  85             90             95
```

Ile	Ala	Ala	Ala	Met	Val	Leu	Arg	Gly	Met	Ala	Lys	Asp	Gly	Gln	Phe
								105					110		

<210> 24  
 <211> 21  
 <212> DNA  
 <213> Artificial Sequence

<220>  
 <223> Description of Artificial Sequence: Synthetic  
 Primer

<400> 24  
 ccagcaaaca acttccccgc c 21

<210> 25  
 <211> 24  
 <212> DNA  
 <213> Artificial Sequence

<220>  
 <223> Description of Artificial Sequence: Synthetic  
 Primer

<400> 25  
 atccttaaac tccgccccat catc 24

<210> 26  
 <211> 28  
 <212> DNA  
 <213> Artificial Sequence

<220>  
 <223> Description of Artificial Sequence: Synthetic  
 Primer

<400> 26  
 gagtgctgtg gagagtgctg ttgatgag 28

<210> 27  
 <211> 27  
 <212> DNA  
 <213> Artificial Sequence

<220>  
 <223> Description of Artificial Sequence: Synthetic  
 Primer

<400> 27  
 ggggataaag gggattgttg atgctgc 27

<210> 28  
<211> 26  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Description of Artificial Sequence: Synthetic  
Primer

<400> 28  
gcaaactgcc catccttagc cattcc 26

<210> 29  
<211> 25  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Description of Artificial Sequence: Synthetic  
Primer

<400> 29  
aaggggattg cgaaggggat aaagg 25

<210> 30  
<211> 27  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Description of Artificial Sequence: Synthetic  
Primer

<400> 30  
ttagcagcaa actttccatc ctagcc 27

<210> 31  
<211> 5897  
<212> DNA  
<213> Borrelia garinii

<400> 31  
cggaatcaa gccacctaaa acaacttccc aaaagtttct caaaaaatat tatattcagc 60  
agtaaattct ataagtcatt aattatttaa tactattcaa cagtaaattc tataagtcatt 120  
taattattta atactattca gcagtaaatt ctataagtca ttaattattt aatactattc 180  
agcagtaaatt tctataagtc attaattatt taatactatt cagcagtaaa ttctataagt 240  
cattaattat ttaatactat tcagcagtaa attctataag tcattaatta tttaatacta 300  
ttcagcagta aattctataa gtcattaatt caattaggta acggattctt agatgtattc 360  
acctcttttg gtggattagt tgcagatgca ttggggttta aagctgatcc aaaaaaatct 420  
gatgtaaaaa cttattttga atctctagct aaaaaattag aagaaacaaa agatgggtta 480  
actaagttgt ccaaaggtaa tgacggtgat actggaaagg ctggtgatgc tgggtgggct 540  
ggtggtggcg ctagtgtgc aggtggcgct ggtgggattg agggcgctat aacagagatt 600  
agcaaatggt tagatgatat ggcaaaagct gctgcggaag ctgcaagtgc tgctactggt 660

aatgcagcaa	ttggggatgt	tgtaaatggt	aatggtggag	cagcaaaaagg	tggtgatgcg	720
gagagtgtta	atgggattgc	taaggggata	aaggggattg	ttgatgctgc	tgagaaggct	780
gatgcgaagg	aagggaaagt	ggatgtggct	ggtgatgctg	gtggggctgg	tggtggcgct	840
ggtgctgcag	gtggcgctgg	tgggattgag	ggcgctataa	cagagattag	caaatggtta	900
gatgatatgg	caaaaagctgc	tgcggttgct	gcaagtgcctg	caagtgcctgc	tactggtaat	960
gcagcaattg	gggatgttgt	taatggtaat	gatggagcag	caaaaaggtgg	tgatgcggcg	1020
agtgttaatg	ggattgctaa	ggggataaaag	gggattgttg	atgctgctga	gaaggctgat	1080
gcgaaggaag	ggaagtggga	tgtggctggt	gatgctgggtg	agggtaacaa	ggatgctggg	1140
aagctgtttg	tgaagaagaa	tgctggtgat	gaggtggtg	aagcaaatga	tgctgggaag	1200
gctgctgctg	cggttgctgc	tgtagtggtg	gagcagatat	taaaagcgat	tgtagtgct	1260
gctgaggggtg	atgataagca	gggtaagaag	gctgcggatg	ctacaaatcc	gattgaggcg	1320
gctattgggg	gtgcggatgc	gggtgctaata	gctgaggcgt	ttaataagat	gaagaaggat	1380
gatcagattg	ctgctgctat	ggttctgagg	ggaatggcta	aggatgggca	gtttgctttg	1440
aaggatgatg	ctgctgctca	tgaagggact	gttaagaatg	ctgttgatat	ggcaaaggcc	1500
gctgcggaag	ctgcaagtgc	tgcaagtgcct	gctactggta	gtacaacgat	tgagatggt	1560
gttaagagtg	gtgaggcaaa	agatggtgat	gcggcgagtg	ttaatgggat	tgctaagggg	1620
ataaagggga	ttgttgatgc	tgctgagaag	gctgatgcga	aggaagggaa	gttgatgtg	1680
gctggtgctg	ctggtacgac	taacgtgaat	gttgggaagt	tgtttgtaa	gaataatggt	1740
aatgaggggtg	gtgatgcaag	tgatgctggg	aaagctgctg	ctgcggttgc	tgctgttagt	1800
ggggagcaga	tattaaaagc	gattgttgat	gctgctaaag	atggtgataa	gcaggggggt	1860
actgatgtaa	aggatgctac	aaatccgatt	gaggcggcta	ttgggggtac	aaatgataat	1920
gatgctgcgg	cgtttgctac	tatgaagaag	gatgatcaga	ttgctgctgc	tatggttctg	1980
aggggaatgg	ctaaggatgg	gcagtttgct	ttgaaggatg	atgctgctaa	ggatggtgat	2040
aaaacggggg	ttgctgcgga	tgctgaaaat	ccgattgacg	cggctattgg	gggtgcggat	2100
gctgatgctg	cggcgtttaa	taaggagggg	atgaagaagg	atgatcagat	tgctgctgct	2160
atggttctga	ggggaatggc	taaggatggg	cagtttgctt	tgacgaataa	tgctgctgct	2220
catgaaggga	ctgttaagaa	tgctgttgat	atggcaaaaag	ctgctgcggt	tgctgaagt	2280
gctgctactg	gcaatgcagc	aattggggat	gttgtaaga	gtaatggtgg	agcagcagca	2340
aaaggtggtg	atgcggcgag	tgtaaatggg	attgctaagg	ggataaaagg	gattgttgat	2400
gctgctgaga	aggctgatgc	gaaggaaagg	aaagtggatg	tggtctggtg	tgctggtgaa	2460
actaacaagg	atgctgggaa	gttgtttggtg	aagaagaatg	gtgatgatgg	tggtgatgca	2520
ggtgatgctg	ggaaggctgc	tgctgcggtt	gctgctgtta	gtggggagca	gatattaaaa	2580
gcgattgttg	atgctgctaa	agatggtgat	aagacggggg	ttactgatgt	aaaggatgct	2640
acaaatccga	ttgacgcggc	tattgggggg	agtgcggatg	ctaattgctga	ggcgtttgat	2700
aagatgaaga	aggatgatca	gattgctgct	gctatggttc	tgaggggaat	ggctaaggat	2760
gggcagtttg	ctttgaagaa	taatgatcat	gataatcata	aggggactgt	taagaatgct	2820
gttgatatgg	caaaggccgc	tgaggaagct	gcaagtgcctg	caagtgcctgc	tactggtaat	2880
gcagcgattg	gggatgttgt	taagaatagt	ggggcagcag	caaaaaggtgg	tgaggcggcg	2940
agtgttaatg	ggattgctaa	ggggataaaag	gggattgttg	atgctgctgg	aaaggctgat	3000
gcgaaggaag	ggaagtggga	tgctactggt	gctgagggtg	cgactaacgt	gaatgctggg	3060
aagttgtttg	tgaagagggc	ggctgatgat	ggtggtgatg	cagatgatgc	tggaaggct	3120
gctgctgcgg	ttgctgcaag	tgctgctact	ggtaatgcag	cgattggaga	tggtgttaat	3180
ggtgatgtgg	caaaaagcaaa	aggtggtgat	gcggcgagtg	ttaatgggat	tgctaagggg	3240
ataaagggga	ttgttgatgc	tgctgagaag	gctgatgcga	aggaagggaa	gttgaatgct	3300
gctggtgctg	aggttacgac	taacgcggat	gctgggaagt	tgtttgtaa	gaatgctggt	3360
aatgtgggtg	gtgaagcagg	tgatgctggg	aaggctgctg	ctgcggttgc	tgctgttagt	3420
ggggagcaga	tattaaaagc	gattgttgat	gctgctaagg	atggtggtga	gaagcagggt	3480
aagaaggctg	cggatgctac	aaatccgatt	gacgcggcta	ttgggggtac	aaatgataat	3540
gatgctgctg	cggcgtttgc	tactatgaag	aaggatgatc	agattgctgc	tgctatggtt	3600
ctgaggggaa	tggctaagga	tgggcaattt	gctttgaagg	atgctgctgc	tgctcatgaa	3660
gggactgtta	agaatgctgt	tgatataata	aaggctgctg	cggaagctgc	aagtgctgca	3720
agtgctgcta	ctggtagtgc	agcaattggg	gatgttggtta	atggtaatgg	agcaacagca	3780
aaaggtggtg	atgcgaagag	tgtaaatggg	attgctaagg	ggataaaagg	gattgttgat	3840
gctgctgaga	aggctgatgc	gaaggaaagg	aagtggatg	tggtctggtga	tgctggtgaa	3900
actaacaagg	atgctgggaa	gttgtttggtg	aagaacaatg	gtaatgaggg	tggtgatgca	3960
gatgatgctg	ggaaggctgc	tgctgcggtt	gctgctgtta	gtggggagca	gatattaaaa	4020
gcgattgttg	atgctgctaa	gggtggtgat	aagacgggta	agaataatgt	gaaggatgct	4080

```

gaaaatccga ttgaggcggc tattgggagt agtgcggatg ctgatgctgc ggcgtttaat 4140
aaggagggga tgaagaagga tgatcagatt gctgctgcta tggttctgag gggaaatggct 4200
aaggatgggc agtttgcttt gacgaatgat gctgctgctc atgaagggaac tgttaagaat 4260
gctgttggga gtgcaacaaa taagaccgtt gttgcttttg ctaacttggg tcgaaagacc 4320
gtgcaagctg ggttgaagaa ggttggggat gttgttaaga atagtgaggc aaaagatggt 4380
gatgcggcga gtgttaatgg gattgctaag gggataaagg ggattgttga tgctgctgag 4440
aaggctgatg cgaaggaagg gaagtggat gtggctgggt ctgctgggtga aactaacaag 4500
gatgctggga agttgtttgt gaagaagaat aatgagggtg gtgaagcaaa tgatgctggg 4560
aaggctgctg ctgcggttgc tgctgttagt ggggagcaga tattaaaagc gattgttgat 4620
gctgctaagg atggtgatga taagcagggt aggatgctac aaatccgatt 4680
gacgcggcta ttgggggtgc aggtgcgggt gctaattgctg ctgcggcggt taataatatg 4740
aagaaggatg atcagattgc tgctgctatg gttctgaggg gaatggctaa ggatgggcag 4800
tttgctttga cgaataatgc tcataactaat cataagggga ctgttaagaa tgctgttgat 4860
atgacaaaag ctgctgcggt tgctgcaagt gctgcaagt ctgctactgg taatgcagca 4920
attggggatg ttgttaatgg taatgatgga gcagcaaaag gtggtgatgc ggcgagtgtt 4980
aatgggattg ctaaggggat aaaggggatt gttgatgctg ctgagaaggc tgatgcgaag 5040
gaagggaagt tgaatgtggc tgggtctgct ggtgctgagg gtaacgaggc tgctgggaag 5100
ctgtttgtga agaagaatgc tgggtgatcat ggtggtgaag caggtgatgc tgggagggct 5160
gctgctgcgg ttgctgctgt tagtggggag cagatattaa aagcgattgt tgatgctgct 5220
aaggatggtg gtgataagca gggtaagaag gctgaggatg ctgaaaatcc gattgacgcg 5280
gctattggga gtacgggtgc ggatgataat gctgctgagg cgtttgctac tatgaagaag 5340
gatgatcaga ttgctgctgc tatggttctg aggggaatgg ctaaggatgg gcagtttgc 5400
ttgaaggatg ctgctcatga taatcataag gggactgtta agaatgctgt tgatataata 5460
aaggctactg cggttgctgc aagtgctgct actggtagta caacgattgg ggatgttgtt 5520
aagaatggtg aggcaaaaagg tggtagggcg aagagtgtta atgggattgc taaggggata 5580
aaggggattg ttgatgctgc tggaaaggct gatgcgaagg aaggggaagt gaatgtggct 5640
ggtgctgctg gtgagggtaa cgaggctgct gggaagctgt ttgtgtaaat tactatagga 5700
ttagaactag tgtacgatat ggtcctttg gttattttgc agctgctaataa gaatttgaag 5760
taagtgaagt taaaattgctg gatgttaatg gaacacattt tattgctaca aaagagaaag 5820
aaatattata tgattcactt gatttaaggg ctctgaggaa aatatttgaa ataacttcaa 5880
agcgaatgtt taagctt 5897

```

<210> 32

<211> 100

<212> PRT

<213> *Borrelia garinii*

<400> 32

```

Val Ile Asn Tyr Leu Ile Leu Phe Ser Ser Lys Phe Tyr Lys Ser Leu
  1                   5                   10                  15

```

```

Ile Gln Leu Gly Asn Gly Phe Leu Asp Val Phe Thr Ser Phe Gly Gly
      20                   25                   30

```

```

Leu Val Ala Asp Ala Leu Gly Phe Lys Ala Asp Pro Lys Lys Ser Asp
      35                   40                   45

```

```

Val Lys Thr Tyr Phe Glu Ser Leu Ala Lys Lys Leu Glu Glu Thr Lys
      50                   55                   60

```

```

Asp Gly Leu Thr Lys Leu Ser Lys Gly Asn Asp Gly Asp Thr Gly Lys
      65                   70                   75                   80

```

```

Ala Gly Asp Ala Gly Gly Ala Gly Gly Gly Ala Ser Ala Ala Gly Gly
      85                   90                   95

```

Ala Gly Gly Ile  
100

<210> 33  
<211> 34  
<212> PRT  
<213> *Borrelia garinii*

<400> 33  
Gly Phe Lys Ala Asp Pro Lys Lys Ser Asp Val Lys Thr Tyr Phe Glu  
1 5 10 15  
Ser Leu Ala Lys Lys Leu Glu Glu Thr Lys Asp Gly Leu Thr Lys Leu  
20 25 30

Ser Lys

<210> 34  
<211> 96  
<212> PRT  
<213> *Borrelia garinii*

<400> 34  
Glu Gly Ala Ile Thr Glu Ile Ser Lys Trp Leu Asp Asp Met Ala Lys  
1 5 10 15  
Ala Ala Ala Glu Ala Ala Ser Ala Ala Thr Gly Asn Ala Ala Ile Gly  
20 25 30  
Asp Val Val Asn Gly Asn Gly Gly Ala Ala Lys Gly Gly Asp Ala Glu  
35 40 45  
Ser Val Asn Gly Ile Ala Lys Gly Ile Lys Gly Ile Val Asp Ala Ala  
50 55 60  
Glu Lys Ala Asp Ala Lys Glu Gly Lys Leu Asp Val Ala Gly Asp Ala  
65 70 75 80  
Gly Gly Ala Gly Gly Gly Ala Gly Ala Ala Gly Gly Ala Gly Gly Ile  
85 90 95

<210> 35  
<211> 198  
<212> PRT  
<213> *Borrelia garinii*

<400> 35  
Glu Gly Ala Ile Thr Glu Ile Ser Lys Trp Leu Asp Asp Met Ala Lys  
1 5 10 15

Ala Ala Ala Val Ala Ala Ser Ala Ala Ser Ala Ala Thr Gly Asn Ala  
                   20                                  25                                  30  
 Ala Ile Gly Asp Val Val Asn Gly Asn Asp Gly Ala Ala Lys Gly Gly  
                   35                                  40                                  45  
 Asp Ala Ala Ser Val Asn Gly Ile Ala Lys Gly Ile Lys Gly Ile Val  
                   50                                  55                                  60  
 Asp Ala Ala Glu Lys Ala Asp Ala Lys Glu Gly Lys Leu Asp Val Ala  
                   65                                  70                                  75                                  80  
 Gly Asp Ala Gly Glu Gly Asn Lys Asp Ala Gly Lys Leu Phe Val Lys  
                                   85                                  90                                  95  
 Lys Asn Ala Gly Asp Glu Gly Gly Glu Ala Asn Asp Ala Gly Lys Ala  
                                   100                                  105                                  110  
 Ala Ala Ala Val Ala Ala Val Ser Gly Glu Gln Ile Leu Lys Ala Ile  
                   115                                  120                                  125  
 Val Asp Ala Ala Glu Gly Asp Asp Lys Gln Gly Lys Lys Ala Ala Asp  
                   130                                  135                                  140  
 Ala Thr Asn Pro Ile Glu Ala Ala Ile Gly Gly Ala Asp Ala Gly Ala  
                   145                                  150                                  155                                  160  
 Asn Ala Glu Ala Phe Asn Lys Met Lys Lys Asp Asp Gln Ile Ala Ala  
                                   165                                  170                                  175  
 Ala Met Val Leu Arg Gly Met Ala Lys Asp Gly Gln Phe Ala Leu Lys  
                   180                                  185                                  190  
 Asp Asp Ala Ala Ala His  
                   195

<210> 36  
 <211> 191  
 <212> PRT  
 <213> *Borrelia garinii*

<400> 36  
 Glu Gly Thr Val Lys Asn Ala Val Asp Met Ala Lys Ala Ala Ala Glu  
                   1                                  5                                  10                                  15  
 Ala Ala Ser Ala Ala Ser Ala Ala Thr Gly Ser Thr Thr Ile Gly Asp  
                   20                                  25                                  30  
 Val Val Lys Ser Gly Glu Ala Lys Asp Gly Asp Ala Ala Ser Val Asn  
                   35                                  40                                  45  
 Gly Ile Ala Lys Gly Ile Lys Gly Ile Val Asp Ala Ala Glu Lys Ala  
                   50                                  55                                  60  
 Asp Ala Lys Glu Gly Lys Leu Asp Val Ala Gly Ala Ala Gly Thr Thr  
                   65                                  70                                  75                                  80



Asn	Val	Asn	Val	Gly	Lys	Leu	Phe	Val	Lys	Asn	Asn	Gly	Asn	Glu	Gly
				85					90					95	
Gly	Asp	Ala	Ser	Asp	Ala	Gly	Lys	Ala	Ala	Ala	Ala	Val	Ala	Ala	Val
			100					105					110		
Ser	Gly	Glu	Gln	Ile	Leu	Lys	Ala	Ile	Val	Asp	Ala	Ala	Lys	Asp	Gly
		115					120					125			
Asp	Lys	Gln	Gly	Val	Thr	Asp	Val	Lys	Asp	Ala	Thr	Asn	Pro	Ile	Glu
	130					135					140				
Ala	Ala	Ile	Gly	Gly	Thr	Asn	Asp	Asn	Asp	Ala	Ala	Ala	Phe	Ala	Thr
145					150					155					160
Met	Lys	Lys	Asp	Asp	Gln	Ile	Ala	Ala	Ala	Met	Val	Leu	Arg	Gly	Met
				165					170					175	
Ala	Lys	Asp	Gly	Gln	Phe	Ala	Leu	Lys	Asp	Asp	Ala	Ala	Lys	Asp	
			180					185					190		

<210> 37

<211> 63

<212> PRT

<213> *Borrelia garinii*

<400> 37

Gly	Asp	Lys	Thr	Gly	Val	Ala	Ala	Asp	Ala	Glu	Asn	Pro	Ile	Asp	Ala
1				5					10					15	
Ala	Ile	Gly	Gly	Ala	Asp	Ala	Asp	Ala	Ala	Ala	Phe	Asn	Lys	Glu	Gly
			20					25					30		
Met	Lys	Lys	Asp	Asp	Gln	Ile	Ala	Ala	Ala	Met	Val	Leu	Arg	Gly	Met
		35					40					45			
Ala	Lys	Asp	Gly	Gln	Phe	Ala	Leu	Thr	Asn	Asn	Ala	Ala	Ala	His	
	50					55					60				

<210> 38

<211> 192

<212> PRT

<213> *Borrelia garinii*

<400> 38

Glu	Gly	Thr	Val	Lys	Asn	Ala	Val	Asp	Met	Ala	Lys	Ala	Ala	Ala	Val
1				5					10					15	
Ala	Ala	Ser	Ala	Ala	Thr	Gly	Asn	Ala	Ala	Ile	Gly	Asp	Val	Val	Lys
			20					25					30		
Ser	Asn	Gly	Gly	Ala	Ala	Ala	Lys	Gly	Gly	Asp	Ala	Ala	Ser	Val	Asn
		35					40					45			

Gly Ile Ala Lys Gly Ile Lys Gly Ile Val Asp Ala Ala Glu Lys Ala  
 50 55 60  
 Asp Ala Lys Glu Gly Lys Leu Asp Val Ala Gly Ala Ala Gly Glu Thr  
 65 70 75 80  
 Asn Lys Asp Ala Gly Lys Leu Phe Val Lys Lys Asn Gly Asp Asp Gly  
 85 90 95  
 Gly Asp Ala Gly Asp Ala Gly Lys Ala Ala Ala Val Ala Ala Val  
 100 105 110  
 Ser Gly Glu Gln Ile Leu Lys Ala Ile Val Asp Ala Ala Lys Asp Gly  
 115 120 125  
 Asp Lys Thr Gly Val Thr Asp Val Lys Asp Ala Thr Asn Pro Ile Asp  
 130 135 140  
 Ala Ala Ile Gly Gly Ser Ala Asp Ala Asn Ala Glu Ala Phe Asp Lys  
 145 150 155 160  
 Met Lys Lys Asp Asp Gln Ile Ala Ala Ala Met Val Leu Arg Gly Met  
 165 170 175  
 Ala Lys Asp Gly Gln Phe Ala Leu Lys Asn Asn Asp His Asp Asn His  
 180 185 190

<210> 39  
 <211> 112  
 <212> PRT  
 <213> *Borrelia garinii*

<400> 39  
 Lys Gly Thr Val Lys Asn Ala Val Asp Met Ala Lys Ala Ala Glu Glu  
 1 5 10 15  
 Ala Ala Ser Ala Ala Ser Ala Ala Thr Gly Asn Ala Ala Ile Gly Asp  
 20 25 30  
 Val Val Lys Asn Ser Gly Ala Ala Ala Lys Gly Gly Glu Ala Ala Ser  
 35 40 45  
 Val Asn Gly Ile Ala Lys Gly Ile Lys Gly Ile Val Asp Ala Ala Gly  
 50 55 60  
 Lys Ala Asp Ala Lys Glu Gly Lys Leu Asp Ala Thr Gly Ala Glu Gly  
 65 70 75 80  
 Thr Thr Asn Val Asn Ala Gly Lys Leu Phe Val Lys Arg Ala Ala Asp  
 85 90 95  
 Asp Gly Gly Asp Ala Asp Asp Ala Gly Lys Ala Ala Ala Val Ala  
 100 105 110

<210> 40  
<211> 174  
<212> PRT  
<213> *Borrelia garinii*

<400> 40  
Ala Ser Ala Ala Thr Gly Asn Ala Ala Ile Gly Asp Val Val Asn Gly  
1 5 10 15  
Asp Val Ala Lys Ala Lys Gly Gly Asp Ala Ala Ser Val Asn Gly Ile  
20 25 30  
Ala Lys Gly Ile Lys Gly Ile Val Asp Ala Ala Glu Lys Ala Asp Ala  
35 40 45  
Lys Glu Gly Lys Leu Asn Ala Ala Gly Ala Glu Gly Thr Thr Asn Ala  
50 55 60  
Asp Ala Gly Lys Leu Phe Val Lys Asn Ala Gly Asn Val Gly Gly Glu  
65 70 75 80  
Ala Gly Asp Ala Gly Lys Ala Ala Ala Ala Val Ala Ala Val Ser Gly  
85 90 95  
Glu Gln Ile Leu Lys Ala Ile Val Asp Ala Ala Lys Asp Gly Gly Glu  
100 105 110  
Lys Gln Gly Lys Lys Ala Ala Asp Ala Thr Asn Pro Ile Asp Ala Ala  
115 120 125  
Ile Gly Gly Thr Asn Asp Asn Asp Ala Ala Ala Ala Phe Ala Thr Met  
130 135 140  
Lys Lys Asp Asp Gln Ile Ala Ala Ala Met Val Leu Arg Gly Met Ala  
145 150 155 160  
Lys Asp Gly Gln Phe Ala Leu Lys Asp Ala Ala Ala Ala His  
165 170

<210> 41  
<211> 195  
<212> PRT  
<213> *Borrelia garinii*

<400> 41  
Glu Gly Thr Val Lys Asn Ala Val Asp Ile Ile Lys Ala Ala Ala Glu  
1 5 10 15  
Ala Ala Ser Ala Ala Ser Ala Ala Thr Gly Ser Ala Ala Ile Gly Asp  
20 25 30

Val Val Asn Gly Asn Gly Ala Thr Ala Lys Gly Gly Asp Ala Lys Ser  
 35 40 45  
 Val Asn Gly Ile Ala Lys Gly Ile Lys Gly Ile Val Asp Ala Ala Glu  
 50 55 60  
 Lys Ala Asp Ala Lys Glu Gly Lys Leu Asp Val Ala Gly Asp Ala Gly  
 65 70 75 80  
 Glu Thr Asn Lys Asp Ala Gly Lys Leu Phe Val Lys Asn Asn Gly Asn  
 85 90 95  
 Glu Gly Gly Asp Ala Asp Asp Ala Gly Lys Ala Ala Ala Ala Val Ala  
 100 105 110  
 Ala Val Ser Gly Glu Gln Ile Leu Lys Ala Ile Val Asp Ala Ala Lys  
 115 120 125  
 Gly Gly Asp Lys Thr Gly Lys Asn Asn Val Lys Asp Ala Glu Asn Pro  
 130 135 140  
 Ile Glu Ala Ala Ile Gly Ser Ser Ala Asp Ala Asp Ala Ala Ala Phe  
 145 150 155 160  
 Asn Lys Glu Gly Met Lys Lys Asp Asp Gln Ile Ala Ala Ala Met Val  
 165 170 175  
 Leu Arg Gly Met Ala Lys Asp Gly Gln Phe Ala Leu Thr Asn Asp Ala  
 180 185 190  
 Ala Ala His  
 195

<210> 42  
 <211> 197  
 <212> PRT  
 <213> *Borrelia garinii*

<400> 42  
 Glu Gly Thr Val Lys Asn Ala Val Gly Ser Ala Thr Asn Lys Thr Val  
 1 5 10 15  
 Val Ala Leu Ala Asn Leu Val Arg Lys Thr Val Gln Ala Gly Leu Lys  
 20 25 30  
 Lys Val Gly Asp Val Val Lys Asn Ser Glu Ala Lys Asp Gly Asp Ala  
 35 40 45  
 Ala Ser Val Asn Gly Ile Ala Lys Gly Ile Lys Gly Ile Val Asp Ala  
 50 55 60  
 Ala Glu Lys Ala Asp Ala Lys Glu Gly Lys Leu Asp Val Ala Gly Ala  
 65 70 75 80  
 Ala Gly Glu Thr Asn Lys Asp Ala Gly Lys Leu Phe Val Lys Lys Asn  
 85 90 95

Asn Glu Gly Gly Glu Ala Asn Asp Ala Gly Lys Ala Ala Ala Val  
                   100                                  105                                  110  
 Ala Ala Val Ser Gly Glu Gln Ile Leu Lys Ala Ile Val Asp Ala Ala  
                   115                                  120                                  125  
 Lys Asp Gly Asp Asp Lys Gln Gly Lys Lys Ala Glu Asp Ala Thr Asn  
                   130                                  135                                  140  
 Pro Ile Asp Ala Ala Ile Gly Gly Ala Gly Ala Gly Ala Asn Ala Ala  
                   145                                  150                                  155                                  160  
 Ala Ala Phe Asn Asn Met Lys Lys Asp Asp Gln Ile Ala Ala Ala Met  
                                   165                                  170                                  175  
 Val Leu Arg Gly Met Ala Lys Asp Gly Gln Phe Ala Leu Thr Asn Asn  
                   180                                  185                                  190  
 Ala His Thr Asn His  
                   195

<210> 43  
 <211> 198  
 <212> PRT  
 <213> *Borrelia garinii*

<400> 43  
 Lys Gly Thr Val Lys Asn Ala Val Asp Met Thr Lys Ala Ala Ala Val  
   1                                  5                                  10                                  15  
 Ala Ala Ser Ala Ala Ser Ala Ala Thr Gly Asn Ala Ala Ile Gly Asp  
                   20                                  25                                  30  
 Val Val Asn Gly Asn Asp Gly Ala Ala Lys Gly Gly Asp Ala Ala Ser  
                   35                                  40                                  45  
 Val Asn Gly Ile Ala Lys Gly Ile Lys Gly Ile Val Asp Ala Ala Glu  
                   50                                  55                                  60  
 Lys Ala Asp Ala Lys Glu Gly Lys Leu Asn Val Ala Gly Ala Ala Gly  
                   65                                  70                                  75                                  80  
 Ala Glu Gly Asn Glu Ala Ala Gly Lys Leu Phe Val Lys Lys Asn Ala  
                                   85                                  90                                  95  
 Gly Asp His Gly Gly Glu Ala Gly Asp Ala Gly Arg Ala Ala Ala Ala  
                   100                                  105                                  110  
 Val Ala Ala Val Ser Gly Glu Gln Ile Leu Lys Ala Ile Val Asp Ala  
                   115                                  120                                  125  
 Ala Lys Asp Gly Gly Asp Lys Gln Gly Lys Lys Ala Glu Asp Ala Glu  
                   130                                  135                                  140  
 Asn Pro Ile Asp Ala Ala Ile Gly Ser Thr Gly Ala Asp Asp Asn Ala

145		150		155		160									
Ala	Glu	Ala	Phe	Ala	Thr	Met	Lys	Lys	Asp	Asp	Gln	Ile	Ala	Ala	Ala
				165					170					175	
Met	Val	Leu	Arg	Gly	Met	Ala	Lys	Asp	Gly	Gln	Phe	Ala	Leu	Lys	Asp
			180					185					190		
Ala	Ala	His	Asp	Asn	His										
		195													

<210> 44  
 <211> 86  
 <212> PRT  
 <213> *Borrelia garinii*

<400> 44
Lys Gly Thr Val Lys Asn Ala Val Asp Ile Ile Lys Ala Thr Ala Val
1 5 10 15
Ala Ala Ser Ala Ala Thr Gly Ser Thr Thr Ile Gly Asp Val Val Lys
20 25 30
Asn Gly Glu Ala Lys Gly Gly Glu Ala Lys Ser Val Asn Gly Ile Ala
35 40 45
Lys Gly Ile Lys Gly Ile Val Asp Ala Ala Gly Lys Ala Asp Ala Lys
50 55 60
Glu Gly Lys Leu Asn Val Ala Gly Ala Ala Gly Glu Gly Asn Glu Ala
65 70 75 80
Ala Gly Lys Leu Phe Val
85

<210> 45  
 <211> 71  
 <212> PRT  
 <213> *Borrelia garinii*

<400> 45
Val Asn Tyr Tyr Arg Ile Arg Thr Ser Val Arg Tyr Glu Ser Phe Gly
1 5 10 15
Tyr Phe Ala Ala Ala Asn Glu Phe Glu Ile Ser Glu Val Lys Ile Ala
20 25 30
Asp Val Asn Gly Thr His Phe Ile Ala Thr Lys Glu Lys Glu Ile Leu
35 40 45
Tyr Asp Ser Leu Asp Leu Arg Ala Arg Gly Lys Ile Phe Glu Ile Thr
50 55 60
Ser Lys Arg Met Phe Lys Leu
65 70

<210> 46  
<211> 8762  
<212> DNA  
<213> *Borrelia afzelii*

<400> 46  
gagagtgcctg ttgatgggggt tagcaagtgg ttagaagaga tgataaaagc tgctaaggag 60  
gctgctacaa aggggtggtac tgggtggtgg agcgaaaaga ttggggatgt tgggtgctgct 120  
aataatcagg gtgctgtagc tgataaggac agtggttaagg ggattgcgaa ggggataaag 180  
gggattgttg atgctgctgg gaaggctttt ggtaaggatg gtaatgcgct gacagggtga 240  
aaagaagttg ctgatgaggg tggggctaac gaggatgcgg ggaagtgtt tgctggtaat 300  
gctggtaatg ctgctgctgc tgacattgcg aaggcggctg gtgctgttac tgcggtagt 360  
ggggagcaga tactgaaagc tattgtttgat ggtgctgggt gtgctggctca agatggtaaa 420  
aaggctgcgg aggcctaagaa tccgattgca gctgcgattg gggctgatgc tgctgggtgcg 480  
gattttggtg atgatatgaa gaagagtgat aagattgctg cggctattgt tttgaggggg 540  
gtggctaaga gtggaaagt ttgctgttgct aatgctgcta agaaggagag tgtgaagagt 600  
gctgtggaga gtgctgttga tgaggtttag aagtggttag aagagatgat aaaagctgct 660  
gggtggggctg ctaagggtgg tactggtgggt aataacgaaa agattgggga ttctgataat 720  
aataagggtg ctgtagctga taaggacagt gttaagggga ttgcgaaggg gataaagggg 780  
attgttgatg ctgctgggaa ggcttttgggt aaggatggta atgcgctgaa ggatgttgca 840  
aaagtgtctg atgatcgccg tggggctaac gcgaatgcag ggaagtgtt tgctggtaat 900  
gctgctgggt gtgccgctga tgctgatgat gctaacattg cgaaggcggc tgggtgctgtt 960  
agtgcggtta gtggggagca gatactgaaa gctattgttg atgctgctgg tgctgctgct 1020  
aatcaggatg gtaagaaggc tgcggatgct aagaatccga ttgcagctgc gattgggact 1080  
aatgatgatg gggcggagtt taaggatgga atgaagaaga gtgataatat tgctgcagct 1140  
attgttttga ggggggtggc taagggtgga aagtttgctg ttgctaattgc tgctaattgat 1200  
agtaaggcga gtgtgaagag tgctgtggag agtgctgttg atgaggttag caagtggtta 1260  
gaagagatga taacagctgc tgggtgaggc gctacaaagg gtggtgatgc tgggtggtgg 1320  
gctgataaga ttggggatgt tgggtgctgct aataatgggt ctgtagctga tgcgagcagt 1380  
gttaaggaga ttgcgaaggg gataaagggg attgttgatg ctgctgggaa ggcttttggc 1440  
aaggatggta atgcgctgaa ggatgttgca gaagtgtctg atgataagaa ggaggcgggg 1500  
aagttgtttg ctggtaatgc tgggtggtgct gttgctgatg ctgctgcgat tgggaaggcg 1560  
gctgggtgctg ttactgcggt tagtggggag cagatactga aagctattgt tgatgctgct 1620  
gggtggtgcgg atcaggcggg taagaaggct gatgcggcta agaatccgat tgcagctgcg 1680  
attggggctg atgctgctgg tgctggtgcg gatttttggt atgatatgaa gaagagaaat 1740  
gataagattg ttgcggctat tgttttgagg ggggtggcta aggatggaaa gtttgctgct 1800  
gctgctaatt atgataatag taaggcgagt gtgaagagt ctgtggagag tgctgttgat 1860  
gaggttagca agtggttaga agagatgata acagctgctg atggggctgc taaagggtgg 1920  
actggtggtg atagcgaaaa gattggggat gctggtgata ataataatgg tgctgtagct 1980  
gatgagaaca gtgttaagga gattgcaaaag gggataaagg ggattgttgc ggctgctggg 2040  
aaggcttttg gcaaggatgg caaggatggg gatgcgctga aggatgttga aacagttgct 2100  
gctgagaatg aggcctaaca ggatgcgggg aagttgtttg ctggtgctaa tggtaatgct 2160  
gggtgctgctg ttggtgacat tgcgaaggcg gctgctgctg ttactgcggt tagtggggag 2220  
cagatactaa aagctattgt tgatgctgct ggtgatgcgg atcaggcggg taagaaggct 2280  
gctgaggcta agaatccgat tgcagctgcg attggggcta atgctgctga taatgcggcg 2340  
gcgttttggt aggatgagat gaagaagagt gataagattg ctgcagctat tgttttgagg 2400  
ggggtggcta aggatggaaa gtttgctgtt gctaattgcta atgatgataa gaaggcgagt 2460  
gtgaagagtg ctgtggagag tgctgtggat gaggttagca agtggttaga agagatgata 2520  
acagctgcta aggaggctgc tacaagggtt ggtactgggt gtaataacga aaagattgga 2580  
gattctgatg ctaataatgg tgcaaggct gatgcgagca gtgttaatgg gattgcgaat 2640  
gggataaagg ggattgttga tgctgctggg aaggcttttg gcaaggagg tagtgcgctg 2700  
aaggatgtta aaacagttgc tgctgagaat gaggctaaca aggatgcggg gaagtgtttt 2760  
gctggtaaga atggtaatgc tgatgctgct gatgctgctg acattgcgaa ggcggctggg 2820  
gctgttagtg cggtttagtg ggagcagata ctgaaagcta ttgttgatgg tgctgggtgat 2880  
gcagctaatac aggcgggtaa aaaggctgct gaggctaaga atccgattgc ggctgcgatt 2940

gggactaatg	aagctggggc	ggagtttgg	gatgatatga	agaagagaaa	tgataagatt	3000
gctgcggcta	ttgttttgag	gggggtggct	aaggatggaa	agtttgctgt	tgctaattgct	3060
gctgctgata	atagtaaggc	gagtgtaag	agtgcgtgtg	atgaggttag	caagtggtta	3120
gaagagatga	taaaggctgc	tggtgaggct	gctacaaaag	gtggtgatgc	tggtggtgg	3180
gctgataaga	ttggggatgc	tggtgataag	ggtgctgtag	ctgatgagc	cagtgttaag	3240
gagattgcga	atgggataaa	ggggattgtt	gatgctgctg	ggaaggcttt	tggcaaggag	3300
ggtagtgcgc	tgaaggatgt	taaaacagtt	gctgctgaga	atgaggctaa	caaggatgcg	3360
gggaagtgt	ttgctggtaa	tgctggtaat	ggtgctgctg	atgacattgc	gaaggcggct	3420
gctgctgtta	ctgcggttag	tggggagcag	atactgaaa	ctattgttga	tgctgctggt	3480
gataaggcta	atcaggatgg	taaaaaggct	gcggtgcta	agaatccgat	tgccgctgcg	3540
attggggctg	ctgatgctgg	tgctgcggcg	gcgtttaatg	agaatgatat	gaagaagagt	3600
gataaagattg	ctgcagctat	tgttttgagg	ggggtggcta	aggatggaaa	gtttgctgct	3660
gctgatgctg	atgctaataa	tagtaaggcg	agcgtgaaga	gtgctgttgg	tgaggtttagc	3720
aagtggttag	aagagatgat	aaaagctgct	ggtgaggctg	caaaagttag	tggtactggt	3780
ggtagcga	agattgggga	tgctgataat	aataagggtg	ctgtagctga	tgcgagcagt	3840
gttaatggga	ttgcgaatgg	gataaagggg	attgtttgatg	ctgctgggaa	ggcttttgg	3900
aaggatggtg	cgctggcagg	tgttgagcgt	gctgctgaga	atgatgataa	gaaggatgcg	3960
gggaagtgt	ttgctggtaa	gaatggtggt	gctggtgctg	ctgatgagc	tgggaaggcg	4020
gctgctgctg	ttactgcggt	tagtggggag	cagatactga	aagctattgt	tgatgctgct	4080
ggtgctgcag	ctaatacaggc	gggtaaaaag	gctgcggatg	ctaagaatcc	gattgcggct	4140
gcgattggga	ctgctgatga	tggggcggag	tttaaggatg	atatgaagaa	gagtgataat	4200
attgctgcgg	ctattgtttt	gaggggggtg	gctaaggatg	gaaagttagc	tggtgcta	4260
gctgatgata	ataaggcag	tgtgaagagt	gctgtggaga	gtgctgttga	tgaggtttagc	4320
aagtggttag	aagagatgat	aacagctgct	ggtgaggctg	caaaagttag	tgctggtggt	4380
ggtgctgata	agattgggga	tgctgcta	aatcagggtg	cgaaggctga	tgagagcagt	4440
gttaatggaa	ttgcaaagg	gataaagggg	attgtttgatg	ctgctgggaa	ggcttttggc	4500
aaggagggtg	gtgcgctgaa	ggatgttgca	aaagtgtgctg	atgatgataa	caaggatgcg	4560
gggaagtgt	ttgctggtaa	tgctggtggt	ggtgctggtg	ctgatattgc	gaaggcggct	4620
gctgctgtta	ctgcggttag	tggggagcag	atactgaaa	ctattgttga	tgctgctggt	4680
gctgcggatc	aggcgggtgc	agctgctggt	gcggctaaga	atccgattgc	ggctgcgatt	4740
ggggctgatg	ctggtgctgc	ggaggagttt	aaggatgaga	tgaagaagag	tgataagatt	4800
gctgcggcta	ttgttttgag	gggggtggct	aagggtggaa	agtttgctgt	tgctgcta	4860
gatgctgcaa	atgtgaagag	tgctgtggag	agtgcgtgtg	gtgaggttag	cgcatggtta	4920
gaagagatga	taacagctgc	tagtgaggct	gctacaaaag	gtggtactgg	tggtactggt	4980
ggtgatagtg	aaaagattgg	ggattctgat	gctaataatg	gtgctgtagc	tgatgcgagc	5040
agtgttaagg	agattgcgaa	ggggataaa	gggattgttg	atgctgctgg	gaaggctttt	5100
ggtaaggatg	gtaatgcgct	gaaggatgtt	gcagaagtgtg	ctgatgatga	ggctaacgcg	5160
gatgcgggga	agttgtttgc	tggtaatgct	ggtaatgctg	ctgctgctga	cgttgcgaag	5220
gcggctggtg	ctgttactgc	ggttagtggtg	gagcagatac	tgaaagctat	tgttgatgct	5280
gctggtgctg	cggatcaggc	gggtgcaaag	gctgatgcgg	ctaagaatcc	gattgcagct	5340
gcgattggga	ctaataagc	tggggcggcg	tttaaggatg	gaatgaagaa	gagaaatgat	5400
aatattgctg	cggctattgt	tttgaggggg	gtggctaaga	gtggaaagt	tgctgttgc	5460
gctgctgatg	ctggtaaaggc	gagagtgtga	agagtgctgt	ggagagtgct	gttgatgagg	5520
ttagcaagtg	gttagaagag	atgataacag	ctgctagtga	ggctgcaaaa	gttggtgctg	5580
tggtgatga	taagattggg	gattctgcta	ataatggtgc	tgtagctgat	gcgggcagtg	5640
ttaagggaat	tgcaagggg	ataaagggga	ttgttgatgc	tgctgggaag	gcttttggtg	5700
aggagggtga	tgcgctgaag	gatgttgcaa	aagttgctga	tgagaatggg	gataacaagg	5760
atgcggggaa	gttgtttgct	ggtgagaatg	gtaatgctgg	tgggtgctgct	gatgctgaca	5820
ttgcgaaggc	ggctgctgct	gttactgcgg	ttagtgggga	gcagatactg	aaagctattg	5880
ttgaggctgc	tgggtgctggt	gatgcagcta	atcaggcggg	taagaaggct	gatgaggcta	5940
agaatccgat	tgcggtcgcg	attgggactg	atgatgctgg	ggcggcgttt	ggtcaggatg	6000
atatgaagaa	gagaaatgat	aatattgctg	cggctattgt	tttgaggggg	gtggctaagg	6060
gtggaaagtt	tgctgttgct	aatgctgcta	atgataagta	ggcgagtgtg	aagagtgctg	6120
tggagagtgc	tggtgatgag	gttagcaagt	ggttagaaga	gataataaca	gctactggga	6180
aggcttttgg	taaggatggt	aatgcgctgg	caggtgttgc	aaaagttagct	gatgatgagg	6240
ctaacgcgga	tgcggggaag	ttgtttgctg	gtgagaatgg	taatgctggt	gctgctgcga	6300
ttgggaaggc	ggctgctgct	gttactgcgg	ttagtgggga	gcagatactg	aaagctattg	6360



```

ttgatgctgc tgggtggtgcg gctcaggtgg gtgctggtgc tgggtgctggct acgaatccga 6420
ttgcagctgc gattggggct gctggtgatg gtgcggatgt tggtaaggat gagatgaaga 6480
agagaaatga taagattgct gctggtatgt ttttgagggg ggtggctaag gatggaaagt 6540
ttgctgctgc tgctaattgat agtaaggcga gtgtgaagag tgctgtggag agtgctgttg 6600
atgaggttag caagtgggta gaagagatga taacagctgc tgatgctgct gctgctaaag 6660
ttggcgatgc tgggtggtggt gctgataaga ttggggatgt tgggtgctgct aataaggggtg 6720
cgaaggctga tgcgagcagt gttaaggaga ttgcgaaggg gataaagggg attggtgatg 6780
ctgctgggaa ggcttttggg ggtgatgctg tgaaggatgt taaagctgct ggtgatgata 6840
acaaggaggc agggaaagtg tttgctgggt ctaatggtaa tgctggtgct aatgctgctg 6900
ctgctgatga cattgcgaag gctggtgctg ctgttagtgc ggtagtggg gagcagatac 6960
tgaaagctat tgtagggcg gctggtgctg cggatcaggc ggggtgtaaag gctgaggagg 7020
ctaagaatcc gattgcagct gcgattggga ctgatgatgc tgggtgctggc gagtttgggtg 7080
agaatgatat gaagaagaat gataatattg ctgcggctat tgttttgagg ggggtgggcta 7140
agagtggaaa gtttctgctc aatgctaatt atgctggtaa gaaggagagt gtgaagagtg 7200
ctgtggatga ggctagcaag tgggtagaag agatgataac agctgctggt gaggctgcta 7260
caaaggggtg tactggtgaa gctagcgaag agattgggga tgttgggtgat aataatcatg 7320
gtgctgtagc tgatgcggac agtggttaagg ggattgcgaa ggggataaag gggattgttg 7380
atgctgctgg gaaggctttt ggtaaggatg gtgcgctgaa ggatgttgca gctgctgctg 7440
gtgatgaggc taacaaggat gcggggaagt tgtttctgctg tcaggatggg ggtggtgctg 7500
atggtgacat tgcgaaggcg gctgctgctg ttactgcggg tagtggggag cagatactga 7560
aagctattgt tgaggctgct ggtgataagg ctaatcaggt ggggtgtaaag gctgctgggtg 7620
cggctacgaa tccgattgca gctgcgattg ggactgatga tgataatgctg gcggcgtttg 7680
ataaggatga gatgaagaag agtaatgata agattgctgc ggctattgtt ttgagggggg 7740
tggctaagga tggaaagttt gctgctaatt ctaatgataa tagtaaggcg agtgtgaaga 7800
gtgctgtgga tgaggttagc aagtggttag aagagatgat aacagctgct agtgatgctg 7860
ctacaaaggg tggtagtggt gaagctagcg aaaagattgg ggattctgat gctaataaag 7920
gtgctggtgc tggggcgggc tttggtgaga atgatatgaa gaagagaaat gataatattg 7980
ctgcagctat tgttttgagg ggggtggcta aggatggaaa gtttctgctt aaggaggatt 8040
attgaactca gctttatagg ggaacagcaa ttcgctagaa aatgattaaa aagcttaact 8100
tcgactgggt cttgccttaa ttttattcct ttgttattat ttatcaatta aattcacttc 8160
ggtttgcttt taaattaatt ctggtatact atgtatacta gatacacaaa ttaaggagaa 8220
gtgaaatgga aaaaatagaa aaatttaaaa acaaatgtca acataaacta caacataaac 8280
taatcgatatt agtatcaaca ctttgctata taaacaataa aaataaaaaa tattcacaaa 8340
gcaacatcct ttattatttt aatgaaaatt taaaaagaaa tgggcaaacc cctattaaaa 8400
taaaaacatt acaaaaactat ctttataaac tggaaaaaga atttgaagta acaactaatt 8460
attataaaca cttggggggtt aattgtggaa ccgaaattta ctataaactt aaatatcaaa 8520
aacaaaaatg ctatcataaa ataaaccaat attttaaaaa gaaaaaagaa attaaattta 8580
acttaagagt aagtgcattt ttaataaaaa aacactcaaa aaaaggaggat gtagaattaa 8640
aggaatgtaa taataataat aataataaag agaaagaaac atcccaaaaa attgaaattt 8700
tacaacaaa agtctatgcc aaaaaatgta aatttttgac aaactactat actaaaattt 8760
ta

```

<210> 47

<211> 202

<212> PRT

<213> Borrelia afzelii

<400> 47

Glu Ser Ala Val Asp Gly Val Ser Lys Trp Leu Glu Glu Met Ile Lys  
1 5 10 15

Ala Ala Lys Glu Ala Ala Thr Lys Gly Gly Thr Gly Gly Gly Ser Glu  
20 25 30

Lys Ile Gly Asp Val Gly Ala Ala Asn Asn Gln Gly Ala Val Ala Asp  
35 40 45

Lys Asp Ser Val Lys Gly Ile Ala Lys Gly Ile Lys Gly Ile Val Asp  
 50 55 60  
 Ala Ala Gly Lys Ala Phe Gly Lys Asp Gly Asn Ala Leu Thr Gly Val  
 65 70 75 80  
 Lys Glu Val Ala Asp Glu Ala Gly Ala Asn Glu Asp Ala Gly Lys Leu  
 85 90 95  
 Phe Ala Gly Asn Ala Gly Asn Ala Ala Ala Asp Ile Ala Lys Ala  
 100 105 110  
 Ala Gly Ala Val Thr Ala Val Ser Gly Glu Gln Ile Leu Lys Ala Ile  
 115 120 125  
 Val Asp Gly Ala Gly Gly Ala Ala Gln Asp Gly Lys Lys Ala Ala Glu  
 130 135 140  
 Ala Lys Asn Pro Ile Ala Ala Ala Ile Gly Ala Asp Ala Ala Gly Ala  
 145 150 155 160  
 Asp Phe Gly Asp Asp Met Lys Lys Ser Asp Lys Ile Ala Ala Ala Ile  
 165 170 175  
 Val Leu Arg Gly Val Ala Lys Ser Gly Lys Phe Ala Val Ala Asn Ala  
 180 185 190  
 Ala Lys Lys Glu Ser Val Lys Ser Ala Val  
 195 200

<210> 48  
 <211> 207  
 <212> PRT  
 <213> *Borrelia afzelii*

<400> 48  
 Glu Ser Ala Val Asp Glu Val Ser Lys Trp Leu Glu Glu Met Ile Lys  
 1 5 10 15  
 Ala Ala Gly Gly Ala Ala Lys Gly Gly Thr Gly Gly Asn Asn Glu Lys  
 20 25 30  
 Ile Gly Asp Ser Asp Asn Asn Lys Gly Ala Val Ala Asp Lys Asp Ser  
 35 40 45  
 Val Lys Gly Ile Ala Lys Gly Ile Lys Gly Ile Val Asp Ala Ala Gly  
 50 55 60  
 Lys Ala Phe Gly Lys Asp Gly Asn Ala Leu Lys Asp Val Ala Lys Val  
 65 70 75 80  
 Ala Asp Asp Ala Ala Gly Ala Asn Ala Asn Ala Gly Lys Leu Phe Ala  
 85 90 95  
 Gly Asn Ala Ala Gly Gly Ala Ala Asp Ala Asp Asp Ala Asn Ile Ala

100					105					110					
Lys	Ala	Ala	Gly	Ala	Val	Ser	Ala	Val	Ser	Gly	Glu	Gln	Ile	Leu	Lys
	115						120					125			
Ala	Ile	Val	Asp	Ala	Ala	Gly	Ala	Ala	Ala	Asn	Gln	Asp	Gly	Lys	Lys
	130					135					140				
Ala	Ala	Asp	Ala	Lys	Asn	Pro	Ile	Ala	Ala	Ala	Ile	Gly	Thr	Asn	Asp
145					150					155					160
Asp	Gly	Ala	Glu	Phe	Lys	Asp	Gly	Met	Lys	Lys	Ser	Asp	Asn	Ile	Ala
				165					170					175	
Ala	Ala	Ile	Val	Leu	Arg	Gly	Val	Ala	Lys	Gly	Gly	Lys	Phe	Ala	Val
			180					185					190		
Ala	Asn	Ala	Ala	Asn	Asp	Ser	Lys	Ala	Ser	Val	Lys	Ser	Ala	Val	
	195						200					205			

<210> 49  
 <211> 210  
 <212> PRT  
 <213> Borrelia afzelii

<400> 49															
Glu	Ser	Ala	Val	Asp	Glu	Val	Ser	Lys	Trp	Leu	Glu	Glu	Met	Ile	Thr
1				5					10					15	
Ala	Ala	Asp	Gly	Ala	Ala	Lys	Gly	Gly	Thr	Gly	Gly	Asn	Ser	Glu	Lys
			20					25					30		
Ile	Gly	Asp	Ala	Gly	Asp	Asn	Asn	Asn	Gly	Ala	Val	Ala	Asp	Glu	Asn
		35					40					45			
Ser	Val	Lys	Glu	Ile	Ala	Lys	Gly	Ile	Lys	Gly	Ile	Val	Ala	Ala	Ala
	50					55					60				
Gly	Lys	Ala	Phe	Gly	Lys	Asp	Gly	Lys	Asp	Gly	Asp	Ala	Leu	Lys	Asp
65					70				75					80	
Val	Glu	Thr	Val	Ala	Ala	Glu	Asn	Glu	Ala	Asn	Lys	Asp	Ala	Gly	Lys
				85				90						95	
Leu	Phe	Ala	Gly	Ala	Asn	Gly	Asn	Ala	Gly	Ala	Ala	Val	Gly	Asp	Ile
			100					105					110		
Ala	Lys	Ala	Ala	Ala	Ala	Val	Thr	Ala	Val	Ser	Gly	Glu	Gln	Ile	Leu
	115						120					125			
Lys	Ala	Ile	Val	Asp	Ala	Ala	Gly	Asp	Ala	Asp	Gln	Ala	Gly	Lys	Lys
	130					135					140				
Ala	Ala	Glu	Ala	Lys	Asn	Pro	Ile	Ala	Ala	Ala	Ile	Gly	Ala	Asn	Ala
145					150					155				160	

Ala Asp Asn Ala Ala Ala Phe Gly Lys Asp Glu Met Lys Lys Ser Asp  
165 170 175

Lys Ile Ala Ala Ala Ile Val Leu Arg Gly Val Ala Lys Asp Gly Lys  
180 185 190

Phe Ala Val Ala Asn Ala Asn Asp Asp Lys Lys Ala Ser Val Lys Ser  
195 200 205

Ala Val  
210

<210> 50

<211> 206

<212> PRT

<213> *Borrelia afzelii*

<400> 50

Glu Ser Ala Val Asp Glu Val Ser Lys Trp Leu Glu Glu Met Ile Thr  
1 5 10 15

Ala Ala Gly Glu Ala Ala Thr Lys Gly Gly Asp Ala Gly Gly Gly Ala  
20 25 30

Asp Lys Ile Gly Asp Val Gly Ala Ala Asn Asn Gly Ala Val Ala Asp  
35 40 45

Ala Ser Ser Val Lys Glu Ile Ala Lys Gly Ile Lys Gly Ile Val Asp  
50 55 60

Ala Ala Gly Lys Ala Phe Gly Lys Asp Gly Asn Ala Leu Lys Asp Val  
65 70 75 80

Ala Glu Val Ala Asp Asp Lys Lys Glu Ala Gly Lys Leu Phe Ala Gly  
85 90 95

Asn Ala Gly Gly Ala Val Ala Asp Ala Ala Ala Ile Gly Lys Ala Ala  
100 105 110

Gly Ala Val Thr Ala Val Ser Gly Glu Gln Ile Leu Lys Ala Ile Val  
115 120 125

Asp Ala Ala Gly Gly Ala Asp Gln Ala Gly Lys Lys Ala Asp Ala Ala  
130 135 140

Lys Asn Pro Ile Ala Ala Ala Ile Gly Ala Asp Ala Ala Gly Ala Gly  
145 150 155 160

Ala Asp Phe Gly Asn Asp Met Lys Lys Arg Asn Asp Lys Ile Val Ala  
165 170 175

Ala Ile Val Leu Arg Gly Val Ala Lys Asp Gly Lys Phe Ala Ala Ala  
180 185 190

Ala Asn Asp Asp Asn Ser Lys Ala Ser Val Lys Ser Ala Val  
195 200 205

<210> 51  
<211> 204  
<212> PRT  
<213> *Borrelia afzelii*

<400> 51

Glu	Ser	Ala	Val	Asp	Glu	Val	Ser	Lys	Trp	Leu	Glu	Glu	Met	Ile	Thr
1				5					10					15	
Ala	Ala	Lys	Glu	Ala	Ala	Thr	Lys	Gly	Gly	Thr	Gly	Gly	Asn	Asn	Glu
			20					25					30		
Lys	Ile	Gly	Asp	Ser	Asp	Ala	Asn	Asn	Gly	Ala	Lys	Ala	Asp	Ala	Ser
		35					40					45			
Ser	Val	Asn	Gly	Ile	Ala	Asn	Gly	Ile	Lys	Gly	Ile	Val	Asp	Ala	Ala
	50					55					60				
Gly	Lys	Ala	Phe	Gly	Lys	Glu	Gly	Ser	Ala	Leu	Lys	Asp	Val	Lys	Thr
65					70					75					80
Val	Ala	Ala	Glu	Asn	Glu	Ala	Asn	Lys	Asp	Ala	Gly	Lys	Leu	Phe	Ala
				85					90					95	
Gly	Lys	Asn	Gly	Asn	Ala	Asp	Ala	Ala	Asp	Ala	Ala	Asp	Ile	Ala	Lys
		100					105						110		
Ala	Ala	Gly	Ala	Val	Ser	Ala	Val	Ser	Gly	Glu	Gln	Ile	Leu	Lys	Ala
		115					120					125			
Ile	Val	Asp	Gly	Ala	Gly	Asp	Ala	Ala	Asn	Gln	Ala	Gly	Lys	Lys	Ala
	130					135					140				
Ala	Glu	Ala	Lys	Asn	Pro	Ile	Ala	Ala	Ala	Ile	Gly	Thr	Asn	Glu	Ala
145				150						155				160	
Gly	Ala	Glu	Phe	Gly	Asp	Asp	Met	Lys	Lys	Arg	Asn	Asp	Lys	Ile	Ala
			165						170					175	
Ala	Ala	Ile	Val	Leu	Arg	Gly	Val	Ala	Lys	Asp	Gly	Lys	Phe	Ala	Val
			180					185					190		
Ala	Asn	Ala	Ala	Ala	Asp	Asn	Ser	Lys	Ala	Ser	Val				
		195					200								

<210> 52  
<211> 203  
<212> PRT  
<213> *Borrelia afzelii*

<400> 52

Glu	Ser	Ala	Val	Asp	Glu	Val	Ser	Lys	Trp	Leu	Glu	Glu	Met	Ile	Lys
1				5					10					15	
Ala	Ala	Gly	Glu	Ala	Ala	Thr	Lys	Gly	Gly	Asp	Ala	Gly	Gly	Gly	Ala

20					25					30					
Asp	Lys	Ile	Gly	Asp	Ala	Gly	Asp	Lys	Gly	Ala	Val	Ala	Asp	Ala	Ser
35				40				45							
Ser	Val	Lys	Glu	Ile	Ala	Asn	Gly	Ile	Lys	Gly	Ile	Val	Asp	Ala	Ala
50				55				60							
Gly	Lys	Ala	Phe	Gly	Lys	Glu	Gly	Ser	Ala	Leu	Lys	Asp	Val	Lys	Thr
65				70				75				80			
Val	Ala	Ala	Glu	Asn	Glu	Ala	Asn	Lys	Asp	Ala	Gly	Lys	Leu	Phe	Ala
				85				90				95			
Gly	Asn	Ala	Gly	Asn	Gly	Ala	Ala	Asp	Asp	Ile	Ala	Lys	Ala	Ala	Ala
100								105				110			
Ala	Val	Thr	Ala	Val	Ser	Gly	Glu	Gln	Ile	Leu	Lys	Ala	Ile	Val	Asp
115				120				125							
Ala	Ala	Gly	Asp	Lys	Ala	Asn	Gln	Asp	Gly	Lys	Lys	Ala	Ala	Asp	Ala
130				135				140							
Lys	Asn	Pro	Ile	Ala	Ala	Ala	Ile	Gly	Ala	Ala	Asp	Ala	Gly	Ala	Ala
145				150				155				160			
Ala	Ala	Phe	Asn	Glu	Asn	Asp	Met	Lys	Lys	Ser	Asp	Lys	Ile	Ala	Ala
				165				170				175			
Ala	Ile	Val	Leu	Arg	Gly	Val	Ala	Lys	Asp	Gly	Lys	Phe	Ala	Ala	Ala
180				185				190							
Asp	Ala	Asp	Ala	Asn	Asn	Ser	Lys	Ala	Ser	Val					
195				200											

<210> 53  
 <211> 200  
 <212> PRT  
 <213> Borrelia afzelii

<400> 53

Glu	Ser	Ala	Val	Gly	Glu	Val	Ser	Lys	Trp	Leu	Glu	Glu	Met	Ile	Lys
1				5				10					15		
Ala	Ala	Gly	Glu	Ala	Ala	Lys	Val	Gly	Gly	Thr	Gly	Gly	Ser	Glu	Lys
20								25				30			
Ile	Gly	Asp	Ala	Asp	Asn	Asn	Lys	Gly	Ala	Val	Ala	Asp	Ala	Ser	Ser
35				40				45							
Val	Asn	Gly	Ile	Ala	Asn	Gly	Ile	Lys	Gly	Ile	Val	Asp	Ala	Ala	Gly
50				55				60							
Lys	Ala	Phe	Gly	Lys	Asp	Gly	Ala	Leu	Ala	Gly	Val	Ala	Ala	Ala	Ala
65				70				75				80			



Ala Ala Ala Ile Gly Ala Asp Ala Gly Ala Ala Glu Glu Phe Lys Asp  
145 150 155 160

Glu Met Lys Lys Ser Asp Lys Ile Ala Ala Ile Val Leu Arg Gly  
165 170 175

Val Ala Lys Gly Gly Lys Phe Ala Val Ala Ala Asn Asp Ala Ala Asn  
180 185 190

Val Lys Ser Ala Val  
195

<210> 55

<211> 199

<212> PRT

<213> *Borrelia afzelii*

<400> 55

Glu Ser Ala Val Gly Glu Val Ser Ala Trp Leu Glu Glu Met Ile Thr  
1 5 10 15

Ala Ala Ser Glu Ala Ala Thr Lys Gly Gly Thr Gly Gly Thr Gly Gly  
20 25 30

Asp Ser Glu Lys Ile Gly Asp Ser Asp Ala Asn Asn Gly Ala Val Ala  
35 40 45

Asp Ala Ser Ser Val Lys Glu Ile Ala Lys Gly Ile Lys Gly Ile Val  
50 55 60

Asp Ala Ala Gly Lys Ala Phe Gly Lys Asp Gly Asn Ala Leu Lys Asp  
65 70 75 80

Val Ala Glu Val Ala Asp Asp Glu Ala Asn Ala Asp Ala Gly Lys Leu  
85 90 95

Phe Ala Gly Asn Ala Gly Asn Ala Ala Ala Asp Val Ala Lys Ala  
100 105 110

Ala Gly Ala Val Thr Ala Val Ser Gly Glu Gln Ile Leu Lys Ala Ile  
115 120 125

Val Asp Ala Ala Gly Ala Ala Asp Gln Ala Gly Ala Lys Ala Asp Ala  
130 135 140

Ala Lys Asn Pro Ile Ala Ala Ala Ile Gly Thr Asn Glu Ala Gly Ala  
145 150 155 160

Ala Phe Lys Asp Gly Met Lys Lys Arg Asn Asp Asn Ile Ala Ala Ala  
165 170 175

Ile Val Leu Arg Gly Val Ala Lys Ser Gly Lys Phe Ala Val Ala Ala  
180 185 190

Ala Asp Ala Gly Lys Ala Arg



195

<210> 56  
<211> 207  
<212> PRT  
<213> *Borrelia afzelii*

<400> 56  
Glu Ser Ala Val Asp Glu Val Ser Lys Trp Leu Glu Glu Met Ile Thr  
1 5 10 15  
Ala Ala Ser Glu Ala Ala Lys Val Gly Ala Gly Gly Asp Asp Lys Ile  
20 25 30  
Gly Asp Ser Ala Asn Asn Gly Ala Val Ala Asp Ala Gly Ser Val Lys  
35 40 45  
Gly Ile Ala Lys Gly Ile Lys Gly Ile Val Asp Ala Ala Gly Lys Ala  
50 55 60  
Phe Gly Lys Glu Gly Asp Ala Leu Lys Asp Val Ala Lys Val Ala Asp  
65 70 75 80  
Glu Asn Gly Asp Asn Lys Asp Ala Gly Lys Leu Phe Ala Gly Glu Asn  
85 90 95  
Gly Asn Ala Gly Gly Ala Ala Asp Ala Asp Ile Ala Lys Ala Ala Ala  
100 105 110  
Ala Val Thr Ala Val Ser Gly Glu Gln Ile Leu Lys Ala Ile Val Glu  
115 120 125  
Ala Ala Gly Ala Gly Asp Ala Ala Asn Gln Ala Gly Lys Lys Ala Asp  
130 135 140  
Glu Ala Lys Asn Pro Ile Ala Ala Ala Ile Gly Thr Asp Asp Ala Gly  
145 150 155 160  
Ala Ala Phe Gly Gln Asp Asp Met Lys Lys Arg Asn Asp Asn Ile Ala  
165 170 175  
Ala Ala Ile Val Leu Arg Gly Val Ala Lys Gly Gly Lys Phe Ala Val  
180 185 190  
Ala Asn Ala Ala Asn Asp Ser Lys Ala Ser Val Lys Ser Ala Val  
195 200 205

<210> 57  
<211> 153  
<212> PRT  
<213> *Borrelia afzelii*

<400> 57  
Glu Ser Ala Val Asp Glu Val Ser Lys Trp Leu Glu Glu Ile Ile Thr  
1 5 10 15

Ala Thr Gly Lys Ala Phe Gly Lys Asp Gly Asn Ala Leu Ala Gly Val  
                   20                                  25                                  30  
 Ala Lys Val Ala Asp Asp Glu Ala Asn Ala Asp Ala Gly Lys Leu Phe  
                   35                                  40                                  45  
 Ala Gly Glu Asn Gly Asn Ala Gly Ala Ala Ala Ile Gly Lys Ala Ala  
                   50                                  55                                  60  
 Ala Ala Val Thr Ala Val Ser Gly Glu Gln Ile Leu Lys Ala Ile Val  
                   65                                  70                                  75                                  80  
 Asp Ala Ala Gly Gly Ala Ala Gln Val Gly Ala Gly Ala Gly Ala Ala  
                                   85                                  90                                  95  
 Thr Asn Pro Ile Ala Ala Ala Ile Gly Ala Ala Gly Asp Gly Ala Asp  
                   100                                  105                                  110  
 Phe Gly Lys Asp Glu Met Lys Lys Arg Asn Asp Lys Ile Ala Ala Ala  
                   115                                  120                                  125  
 Ile Val Leu Arg Gly Val Ala Lys Asp Gly Lys Phe Ala Ala Ala Ala  
                   130                                  135                                  140  
 Asn Asp Ser Lys Ala Ser Val Lys Ser  
                   145                                  150

<210> 58  
 <211> 202  
 <212> PRT  
 <213> *Borrelia afzelii*

<400> 58  
 Glu Ser Ala Val Asp Glu Val Ser Lys Trp Leu Glu Glu Met Ile Thr  
           1                                  5                                  10                                  15  
 Ala Ala Asp Ala Ala Ala Ala Lys Val Gly Asp Ala Gly Gly Gly Ala  
                   20                                  25                                  30  
 Asp Lys Ile Gly Asp Val Gly Ala Ala Asn Lys Gly Ala Lys Ala Asp  
                   35                                  40                                  45  
 Ala Ser Ser Val Lys Glu Ile Ala Lys Gly Ile Lys Gly Ile Val Asp  
                   50                                  55                                  60  
 Ala Ala Gly Lys Ala Phe Gly Gly Asp Ala Leu Lys Asp Val Lys Ala  
                   65                                  70                                  75                                  80  
 Ala Gly Asp Asp Asn Lys Glu Ala Gly Lys Leu Phe Ala Gly Ala Asn  
                                   85                                  90                                  95  
 Gly Asn Ala Gly Ala Asn Ala Ala Ala Ala Asp Asp Ile Ala Lys Ala  
                   100                                  105                                  110  
 Ala Gly Ala Val Ser Ala Val Ser Gly Glu Gln Ile Leu Lys Ala Ile

115	120	125
Val Glu Ala Ala Gly Ala Ala Asp Gln Ala Gly Val Lys Ala Glu Glu		
130	135	140
Ala Lys Asn Pro Ile Ala Ala Ala Ile Gly Thr Asp Asp Ala Gly Ala		
145	150	155 160
Ala Glu Phe Gly Glu Asn Asp Met Lys Lys Asn Asp Asn Ile Ala Ala		
	165	170 175
Ala Ile Val Leu Arg Gly Val Ala Lys Ser Gly Lys Phe Ala Ala Asn		
	180	185 190
Ala Asn Asp Ala Gly Lys Lys Glu Ser Val		
195	200	

<210> 59  
 <211> 201  
 <212> PRT  
 <213> *Borrelia afzelii*

<400> 59

Lys Ser Ala Val Asp Glu Ala Ser Lys Trp Leu Glu Glu Met Ile Thr		
1	5	10 15
Ala Ala Gly Glu Ala Ala Thr Lys Gly Gly Thr Gly Glu Ala Ser Glu		
	20	25 30
Lys Ile Gly Asp Val Gly Asp Asn Asn His Gly Ala Val Ala Asp Ala		
	35	40 45
Asp Ser Val Lys Gly Ile Ala Lys Gly Ile Lys Gly Ile Val Asp Ala		
	50	55 60
Ala Gly Lys Ala Phe Gly Lys Asp Gly Ala Leu Lys Asp Val Ala Ala		
	65	70 75 80
Ala Ala Gly Asp Glu Ala Asn Lys Asp Ala Gly Lys Leu Phe Ala Gly		
	85	90 95
Gln Asp Gly Gly Gly Ala Asp Gly Asp Ile Ala Lys Ala Ala Ala Ala		
	100	105 110
Val Thr Ala Val Ser Gly Glu Gln Ile Leu Lys Ala Ile Val Glu Ala		
	115	120 125
Ala Gly Asp Lys Ala Asn Gln Val Gly Val Lys Ala Ala Gly Ala Ala		
	130	135 140
Thr Asn Pro Ile Ala Ala Ala Ile Gly Thr Asp Asp Asp Asn Ala Ala		
	145	150 155 160
Ala Phe Asp Lys Asp Glu Met Lys Lys Ser Asn Asp Lys Ile Ala Ala		
	165	170 175

Ala Ile Val Leu Arg Gly Val Ala Lys Asp Gly Lys Phe Ala Ala Asn  
180 185 190

Ala Asn Asp Asn Ser Lys Ala Ser Val  
195 200

<210> 60

<211> 82

<212> PRT

<213> Borrelia afzelii

<400> 60

Lys Ser Ala Val Asp Glu Val Ser Lys Trp Leu Glu Glu Met Ile Thr  
1 5 10 15

Ala Ala Ser Asp Ala Ala Thr Lys Gly Gly Thr Gly Glu Ala Ser Glu  
20 25 30

Lys Ile Gly Asp Ser Asp Ala Asn Lys Gly Ala Gly Ala Gly Ala Ala  
35 40 45

Phe Gly Glu Asn Asp Met Lys Lys Arg Asn Asp Asn Ile Ala Ala Ala  
50 55 60

Ile Val Leu Arg Gly Val Ala Lys Asp Gly Lys Phe Ala Val Lys Glu  
65 70 75 80

Asp Tyr

<210> 61

<211> 179

<212> PRT

<213> Borrelia afzelii

<400> 61

Met Glu Lys Ile Glu Lys Phe Lys Asn Lys Cys Gln His Lys Leu Gln  
1 5 10 15

His Lys Leu Ile Val Leu Val Ser Thr Leu Cys Tyr Ile Asn Asn Lys  
20 25 30

Asn Lys Lys Tyr Ser Gln Ser Asn Ile Leu Tyr Tyr Phe Asn Glu Asn  
35 40 45

Leu Lys Arg Asn Gly Gln Thr Pro Ile Lys Ile Lys Thr Leu Gln Asn  
50 55 60

Tyr Leu Tyr Lys Leu Glu Lys Glu Phe Glu Val Thr Thr Asn Tyr Tyr  
65 70 75 80

Lys His Leu Gly Val Asn Cys Gly Thr Glu Ile Tyr Tyr Lys Leu Lys  
85 90 95

Tyr Gln Lys Gln Lys Cys Tyr His Lys Ile Asn Gln Tyr Phe Lys Lys

100

105

110

Lys Lys Glu Ile Lys Phe Asn Leu Arg Val Ser Ala Phe Phe Asn Lys  
 115 120 125

Lys His Ser Lys Lys Gly Ser Val Glu Leu Lys Glu Cys Asn Asn Asn  
 130 135 140

Asn Asn Asn Lys Glu Lys Glu Thr Ser Gln Lys Ile Glu Ile Leu Gln  
 145 150 155 160

Thr Lys Val Tyr Ala Lys Lys Cys Lys Phe Leu Thr Asn Tyr Tyr Thr  
 165 170 175

Lys Ile Leu

<210> 62

<211> 2775

<212> DNA

<213> *Borrelia garinii*

<400> 62

cggaaatcaa gccacctaata acaacttccc aaaagtttct caaaaaatat tatattcagc 60  
 agtaaatctc ataagtcatt aattatttaa tactattcaa cagtaaattc tataagtcac 120  
 taattattta atactattca gcagtaaatt ctataagtc ttaattattt aatactattc 180  
 agcagtaaatt tctataagtc attaattatt taatactatt cagcagtaaa ttctataagt 240  
 cattaattat ttaatactat tcagcagtaa attctataag tcattaatta tttaatacta 300  
 ttcagcagta aattctataa gtcattaatt caattaggta acggattctt agatgtattc 360  
 acctcttttg gtggattagt tgcagatgca ttggggttta aagctgatcc aaaaaaatct 420  
 gatgtaaaaa cttattttga atctctagct aaaaaattag aagaaacaaa agatggttta 480  
 actaagttgt ccaaaggtaa tgacgggtgat actggaaagg ctgggtgatgc tgggtggggct 540  
 ggtggtggcg ctagtgtctc aggtggcgct ggtgggattg agggcgctat aacagagatt 600  
 agcaaatggt tagatgatat ggcaaaagct gctgcggaag ctgcaagtgc tgctactggt 660  
 aatgcagcaa ttggggatgt tgttaatggt aatgggtggag cagcaaaagg tgggtgatgcg 720  
 gagagtgtta atgggattgc taaggggata aaggggattg ttgatgctgc tgagaaggct 780  
 gatgcgaagg aagggaaagt ggatgtggct ggtgatgctg gtggggctgg tgggtggcgct 840  
 ggtgctgcag gtggcgctgg tgggattgag ggcgctataa cagagattag caaatgggta 900  
 gatgatattg caaaagctgc tgcggttgct gcaagtgctg caagtgctgc tactggtaac 960  
 gcagcaattg gggatgttgt taatggtaac gatggagcag caaaagggtg tgatgcggcg 1020  
 agtggttaatg ggattgctaa ggggataaag gggattgttg atgctgctga gaaggctgat 1080  
 gcgaagggaag ggaagtggga tgtggctggt gatgctgggt agggtaacaa ggatgctggg 1140  
 aagctgtttg tgaagaagaa tgctggtgat gaggggtggt aagcaaatga tgctgggaag 1200  
 gctgctgctg cggttgctgc tgttagtggg gagcagatat taaaagcgat tgttgatgct 1260  
 gctgaggggt atgataagca gggtaagaag gctgcggatg ctacaaatcc gattgaggcg 1320  
 gctattgggg gtgcggatgc ggggtgctaatt gctgaggcgt ttaataagat gaagaaggat 1380  
 gatcagattg ctgctgctat ggttctgagg ggaatggcta aggatgggca gtttgctttg 1440  
 aaggatgatg ctgctgctca tgaagggaact gttaagaatg ctggtgatat ggcaaaggcc 1500  
 gctgcggaag ctgcaagtgc tgcaagtgct gctactggta gtacaacgat tggagatgtt 1560  
 gttaagagtg gtgaggcaaa agatggtgat gcggcgagtg ttaatgggat tgctaagggg 1620  
 ataaagggga ttgttgatgc tgctgagaag gctgatgcga aggaagggaa gttggatgtg 1680  
 gctggtgctg ctggtacgac taacgtgaat gttgggaagt tgtttgtgaa gaataatggt 1740  
 aatgaggggt gtgatgcaag tgatgctggg aaagctgctg ctgcggttgc tgctgttagt 1800  
 ggggagcaga tattaagagc gattgttgat gctgctaaag atggtgataa gcaggggggt 1860  
 actgatgtaa aggatgctac aaatccgatt gaggcggcta ttgggggtac aaatgataat 1920  
 gatgctgcgg cgtttgctac tatgaagaag gatgatcaga ttgctgctgc tatgggttctg 1980

aggggaatgg	ctaaggatgg	gcagtttgct	ttgaaggatg	atgctgctaa	ggatggatgat	2040
aaaacggggg	ttgctgcgga	tgctgaaaat	ccgattgacg	cggctattgg	gggtgcggat	2100
gctgatgctg	cggcggttaa	taaggagggg	atgaagaagg	atgatcagat	tgctgctgct	2160
atggttctga	ggggaatggc	taaggatggg	cagtttgctt	tgacgaataa	tgctgctgct	2220
catgaaggga	ctgttaagaa	tgctggtgat	atggcaaaag	ctgctgcggt	tgctgcaagt	2280
gctgctactg	gcaatgcagc	aattggggat	gttgtaaga	gtaatggtgg	agcagcagca	2340
aaaggtggtg	atgcggcgag	tgtaaatggg	attgctaagg	ggataaaggg	gattgttgat	2400
gctgctgaga	aggctgatgc	gaagggaagg	aagttggatg	tggctggtgc	tgctggtgaa	2460
actaacaagg	atgctgggaa	gttgtttggt	aagaagaatg	gtgatgatgg	tggtgatgca	2520
ggtgatgctg	ggaaggctgc	tgctgcgggt	gctgctgtta	gtggggagca	gatattaaaa	2580
gcgattgttg	atgctgctaa	agatgggtgat	aagacggggg	ttactgatgt	aaaggatgct	2640
acaaatccga	ttgacgcggc	tattgggggg	agtgcggatg	ctaatactga	ggcggttgat	2700
aagatgaaga	aggatgatca	gattgctgct	gctatggttc	tgaggggaat	ggctaaggat	2760
gggcagtttg	ctttg					2775

<210> 63

<211> 2075

<212> DNA

<213> *Borrelia garinii*

<400> 63

ataaagggga	ttgttgatgc	tgctgagaag	gctgatgcga	aggaagggaa	gttggatgtg	60
gctggtgatg	ctggtgaaac	taacaaggat	gctgggaagt	tgtttgtaaa	gaacaatggt	120
aatgaggggtg	gtgatgcaga	tgatgctggg	aaggctgctg	ctgcggttgc	tgctgttagt	180
ggggagcaga	tattaaaagc	gattgttgat	gctgctaagg	gtggtgataa	gacgggtaag	240
aataatgtga	aggatgctga	aaatccgatt	gaggcggcta	ttgggagtag	tgcggtatgct	300
gatgctgcgg	cgtttaataa	ggaggggatg	aagaaggatg	atcagattgc	tgctgctatg	360
gttctgaggg	gaatggctaa	ggatgggcag	tttgctttga	cgaatgatgc	tgctgctcat	420
gaagggactg	ttaagaatgc	tggtgggagt	gcaacaaata	agaccgttgt	tgctttggct	480
aacttggttc	gaaagaccgt	gcaagctggg	ttgaagaagg	ttggggatgt	tgtaagaat	540
agtgaggcaa	aagatggtga	tgcggcgagt	gttaatggga	ttgctaaggg	gataaagggg	600
attgttgatg	ctgctgagaa	ggctgatgct	aaggaaggga	agttggatgt	ggctggtgct	660
gctggtgaaa	ctaacaagga	tgctgggaag	ttgtttgtga	agaagaataa	tgaggggtgt	720
gaagcaaatg	atgctgggaa	ggctgctgct	gcggttgctg	ctgttagtgg	ggagcagata	780
ttaaaagcga	ttgttgatgc	tgctaaggat	ggtgatgata	agcagggtaa	gaaggctgag	840
gatgctacaa	atccgattga	cgcggctatt	gggggtgcag	gtgcgggtgc	taatgctgct	900
gcggcggttta	ataatatgaa	gaaggatgat	cagattgctg	ctgctatggt	tctgagggga	960
atggctaagg	atgggcagtt	tgctttgacg	aataatgctc	ataactaatca	taaggggact	1020
gttaagaatg	ctgttgatat	gacaaaagct	gctgcggttg	ctgcaagtgc	tgcaagtgct	1080
gctactggta	atgcagcaat	tggggatgtt	gttaatggta	atgatggagc	agcaaaagggt	1140
ggtgatgcgg	cgagtgttaa	tgggattgct	aaggggataa	aggggattgt	tgatgctgct	1200
gagaaggctg	atgcgaagga	agggaaagttg	aatgtggctg	gtgctgctgg	tgctgaggggt	1260
aacgaggctg	ctgggaagct	gtttgtgaag	aagaatgctg	gtgatcatgg	tggtgaagca	1320
ggtgatgctg	ggagggctgc	tgctgcgggt	gctgctgtta	gtggggagca	gatattaaaa	1380
gcgattgttg	atgctgctaa	ggatgggtgt	gataagcagg	gtaagaaggc	tgaggatgct	1440
gaaaatccga	ttgacgcggc	tattgggagt	acgggtgcgg	atgataatgc	tgctgaggcg	1500
tttgctacta	tgaagaagga	tgatcagatt	gctgctgcta	tggttctgag	gggaatggct	1560
aaggatgggc	agtttgcttt	gaaggatgct	gctcatgata	atcataaggg	gactgttaag	1620
aatgctgttg	atataataaa	ggctactgct	gttgctgcaa	gtgctgctac	tggtagtaca	1680
acgattgggg	atgttggttaa	gaatggtgag	gcaaaagggtg	gtgaggcgaa	gagtgttaaat	1740
gggattgcta	aggggataaa	ggggattgtt	gatgctgctg	gaaaggctga	tgcaaggaa	1800
gggaagttga	atgtggctgg	tgctgctggt	gagggtaacg	aggctgctgg	gaagctgttt	1860
gtgtaaatta	ctataggatt	agaactagtg	tacgatatga	gtcctttggt	tattttgcag	1920
ctgctaataga	atgtgaaata	agtgaagtta	aaattgcgga	tgtaaatgga	acacatttta	1980
ttgctacaaa	agagaaagaa	atattatatg	attcacttga	tttaagggct	cgtggaaaaa	2040
tatttgaaat	aacttcaaag	cgaatgttta	agctt			2075

<210> 64  
 <211> 2775  
 <212> DNA  
 <213> *Borrelia garinii*

<400> 64  
 cggaatcaa gccacctaaa acaacttccc aaaagtttct caaaaaatat tatattcagc 60  
 agtaaattct ataagtcatt aattatttaa tactattcaa cagtaaattc tataagtcac 120  
 taattattta atactattca gcagtaaatt ctataagtca ttaattattt aatactattc 180  
 agcagtaaatt tctataagtc attaattatt taatactatt cagcagtaaa ttctataagt 240  
 cattaattat ttaatactat tcagcagtaa attctataag tcattaatta tttaatacta 300  
 ttcagcagta aattctataa gtcattaatt caattaggta acggattctt agatgtattc 360  
 acctcttttg gtggattagt tgcagatgca ttggggttta aagctgatcc aaaaaaatct 420  
 gatgtaaaaa cttattttga atctctagct aaaaaattag aagaaacaaa agatggttta 480  
 actaagttgt ccaaaggtaa tgacggtgat actggaaagg ctggtgatgc tgggtggggct 540  
 ggtggtggcg ctagtgtgct aggtggcgct ggtgggattg agggcgctat aacagagatt 600  
 agcaaaggt tagatgatat ggcaaaagct gctgcggaag ctgcaagtgc tgctactggt 660  
 aatgcagcaa ttggggatgt tgttaatggt aatggtggag cagcaaaagg tggatgatgcg 720  
 gagagtgtta atgggattgc taaggggata aaggggattg ttgatgctgc tgagaaggct 780  
 gatgcgaagg aagggaagtt ggatgtggct ggtgatgctg gtggggctgg tgggtggcgct 840  
 ggtgctgcag gtggcgctgg tgggattgag ggcgctataa cagagattag caaatgggta 900  
 gatgatattg caaaagctgc tgcggttgct gcaagtgcct caagtgcctc tactggtaac 960  
 gcagcaattg gggatgttgt taatggtaac gatggagcag caaaagggtg tgatgcggcg 1020  
 agtggttaag ggattgctaa ggggataaag gggattgttg atgctgctga gaaggctgat 1080  
 gcgaaggaa ggaagtggga tgtggctggg gatgctgggt agggtaacaa ggatgctggg 1140  
 aagctgtttg tgaagaagaa tgctggtgat gagggtgggt aagcaaatga tgctgggaag 1200  
 gctgctgctg cggttgctgc tgttagtggg gacagatat taaaagcgat tgttgatgct 1260  
 gctgaggggt atgataagca gggtaagaag gctgcggatg ctacaaatcc gattgaggcg 1320  
 gctattgggg gtgcggatgc ggggtgcta gctgaggcgt ttaataagat gaagaaggat 1380  
 gatcagattg ctgctgctat ggttctgagg ggaatggcta aggatgggca gtttgctttg 1440  
 aaggatgatg ctgctgctca tgaagggaact gttaagaatg ctgttgatat ggcaaaggcc 1500  
 gctgcggaag ctgcaagtgc tgcaagtgc gctactggta gtacaacgat tggagatgtt 1560  
 gttaagagtg gtgaggcaaa agatggtgat gcggcgagt ttaatgggat tgctaagggg 1620  
 ataaagggga ttgttgatgc tgctgagaag gctgatgcga aggaaggga gttggatgtg 1680  
 gctggtgctg ctggtacgac taacgtgaat gttgggaagt tgtttgtgaa gaataatggt 1740  
 aatgaggggt gtgatgcaag tgatgctggg aaagctgctg ctgcggttgc tgctgttagt 1800  
 ggggagcaga tattaanaagc gattgttgat gctgctaaag atggtgataa gcaggggggt 1860  
 actgatgtaa aggatgctac aaatccgatt gaggcggcta ttgggggtac aaatgataat 1920  
 gatgctgcgg cgtttgctac tatgaagaag gatgatcaga ttgctgctgc tatggttctg 1980  
 aggggaatgg ctaaggatgg gcagtttgct ttgaaggatg atgctgctaa ggatgggtgat 2040  
 aaaacggggg ttgctgcgga tgctgaaaat ccgattgacg cggctatttg ggggtcggat 2100  
 gctgatgctg cggcgtttaa taaggagggg atgaagaagg atgatcagat tgctgctgct 2160  
 atggttctga ggggaatggc taaggatggg cagtttgctt tgacgaataa tgctgctgct 2220  
 catgaaggga ctgttaagaa tgctgttgat atggcaaaag ctgctgcggg tgctgcaagt 2280  
 gctgctactg gcaatgcagc aattggggat gttgttaaga gtaatgggtg agcagcagca 2340  
 aaagggtggt atgcggcgag tgttaatggg attgctaagg ggataaaggg gattgttgat 2400  
 gctgctgaga aggctgatgc gaagggaagg aagttggatg tggctgggtg tgctggtgaa 2460  
 actaacaagg atgctgggaa gttgtttgtg aagaagaatg gtgatgatgg tggatgatga 2520  
 ggtgatgctg ggaaggctgc tgctgcgggt gctgctgtta gtggggagca gatattaaaa 2580  
 gcgattgttg atgctgctaa agatggtgat aagacggggg ttactgatgt aaaggatgct 2640  
 acaaatccga ttgacgcggc tattgggggg agtgcggtat ctaatgctga ggcgtttgat 2700  
 aagatgaaga aggatgatca gattgctgct gctatggttc tgaggggaat ggctaaggat 2760  
 gggcagtttg ctttg 2775

<210> 65  
 <211> 2075  
 <212> DNA  
 <213> *Borrelia garinii*

<400> 65  
 ataaagggga ttgttgatgc tgctgagaag gctgatgcga aggaagggaa gttggatgtg 60  
 gctggtgatg ctggtgaaac taacaaggat gctgggaagt tgtttgtgaa gaacaatggt 120  
 aatgaggggtg gtgatgcaga tgatgctggg aaggctgctg ctgcggttgc tgctgttagt 180  
 ggggagcaga tattaaaagc gattgttgat gctgctaagg gtggtgataa gacgggtaag 240  
 aataatgtga aggatgctga aaatccgatt gaggcggcta ttgggagtag tgcggatgct 300  
 gatgctgcgg cgtttaataa ggaggggatg aagaaggatg atcagattgc tgctgctatg 360  
 gttctgaggg gaatggctaa ggatgggcag tttgctttga cgaatgatgc tgctgctcat 420  
 gaagggactg ttaagaatgc tgttgggagt gcaacaaata agaccgttgt tgctttggct 480  
 aacttggttc gaaagaccgt gcaagctggg ttgaagaagg ttggggatgt tgtaagaat 540  
 agtgaggcaa aagatggtga tgcggcgagt gttaatggga ttgctaaggg gataaagggg 600  
 attgttgatg ctgctgagaa ggctgatgcg aaggaaggga agttggatgt ggctgggtgct 660  
 gctggtgaaa ctaacaagga tgctgggaag ttgtttgtga agaagaataa tgaggggtggt 720  
 gaagcaaatt atgctgggaa ggctgctgct gcggttgctg ctgttagtgg ggagcagata 780  
 ttaaaagcga ttgttgatgc tgctaaggat ggtgatgata agcagggtaa gaaggctgag 840  
 gatgctacaa atccgattga cgcggctatt gggggtgcag gtgcgggtgc taatgctgct 900  
 gcggcgttta ataatatgaa gaaggatgat cagattgctg ctgctatggt tctgagggga 960  
 atggctaagg atgggcagtt tgctttgacg aataatgctc atactaatca taaggggact 1020  
 gttaagaatg ctggtgatat gacaaaagct gctgcggttg ctgcaagtgc tgcaagtgct 1080  
 gctactggta atgcagcaat tggggatggt gttaatggta atgatggagc agcaaaaggt 1140  
 ggtgatgcgg cgagtgttaa tgggattgct aaggggataa aggggattgt tgatgctgct 1200  
 gagaaggctg atgcgaagga agggaagttg aatgtggctg gtgctgctgg tgctgagggg 1260  
 aacgaggctg ctgggaagct gtttgtgaag aagaatgctg gtgatcatgg tggatgaagca 1320  
 ggtgatgctg ggagggctgc tgctgcggtt gctgctgtta gtggggagca gatattaaaa 1380  
 gcgattgttg atgctgctaa ggatggtggt gataagcagg gtaagaaggc tgaggatgct 1440  
 gaaaatccga ttgacgcggc tattgggagt acgggtgcgg atgataatgc tgctgaggcg 1500  
 tttgctacta tgaagaagga tgatcagatt gctgctgcta tggttctgag gggaatggct 1560  
 aaggatgggc agtttgcttt gaaggatgct gctcatgata atcataaggg gactgttaag 1620  
 aatgctgttg atataataaa ggctactgcg gttgctgcaa gtgctgctac tggtagtaca 1680  
 acgattgggg atgttgtaa gaatggtgag gcaaaagggt gtgaggcgaa gagtgttaat 1740  
 gggattgcta aggggataaa ggggattggt gatgctgctg gaaaggctga tgcgaaggaa 1800  
 ggggaagtga atgtggctgg tgctgctggt gagggtaacg aggctgctgg gaagctgttt 1860  
 gtgtaaatta ctataggatt agaactagtg tacgatatga gtcctttggt tattttgcag 1920  
 ctgctaataa atttgaataa agtgaagtta aaattgcgga tgttaatgga acacatttta 1980  
 ttgctacaaa agagaaagaa atattatatg attcacttga ttttaagggt cgtggaaaaa 2040  
 tatttgaaat aacttcaaag cgaatgttta agctt 2075

<210> 66  
 <211> 184  
 <212> PRT  
 <213> *Borrelia garinii*

<400> 66  
 Glu Gly Thr Val Lys Asn Ala Val Asp Met Ala Lys Ala Ala Val  
 1 5 10 15  
 Ala Ala Ser Ala Ala Thr Gly Asn Ala Ala Ile Gly Asp Val Val Lys  
 20 25 30  
 Ser Asn Gly Gly Ala Ala Ala Lys Gly Gly Asp Ala Ala Ser Val Asn  
 35 40 45



Gly Ile Ala Lys Gly Ile Lys Gly Ile Val Asp Ala Ala Glu Lys Ala  
 50 55 60  
 Asp Ala Lys Glu Gly Lys Leu Asp Val Ala Gly Ala Ala Gly Glu Thr  
 65 70 75 80  
 Asn Lys Asp Ala Gly Lys Leu Phe Val Lys Lys Asn Gly Asp Asp Gly  
 85 90 95  
 Gly Asp Ala Gly Asp Ala Gly Lys Ala Ala Ala Val Ala Ala Val  
 100 105 110  
 Ser Gly Glu Gln Ile Leu Lys Ala Ile Val Asp Ala Ala Lys Asp Gly  
 115 120 125  
 Asp Lys Thr Gly Val Thr Asp Val Lys Asp Ala Thr Asn Pro Ile Asp  
 130 135 140  
 Ala Ala Ile Gly Gly Ser Ala Asp Ala Asn Ala Glu Ala Phe Asp Lys  
 145 150 155 160  
 Met Lys Lys Asp Asp Gln Ile Ala Ala Ala Met Val Leu Arg Gly Met  
 165 170 175  
 Ala Lys Asp Gly Gln Phe Ala Leu  
 180

<210> 67  
 <211> 140  
 <212> PRT  
 <213> *Borrelia garinii*

<400> 67  
 Ile Lys Gly Ile Val Asp Ala Ala Glu Lys Ala Asp Ala Lys Glu Gly  
 1 5 10 15  
 Lys Leu Asp Val Ala Gly Asp Ala Gly Glu Thr Asn Lys Asp Ala Gly  
 20 25 30  
 Lys Leu Phe Val Lys Asn Asn Gly Asn Glu Gly Gly Asp Ala Asp Asp  
 35 40 45  
 Ala Gly Lys Ala Ala Ala Ala Val Ala Ala Val Ser Gly Glu Gln Ile  
 50 55 60  
 Leu Lys Ala Ile Val Asp Ala Ala Lys Gly Gly Asp Lys Thr Gly Lys  
 65 70 75 80  
 Asn Asn Val Lys Asp Ala Glu Asn Pro Ile Glu Ala Ala Ile Gly Ser  
 85 90 95  
 Ser Ala Asp Ala Asp Ala Ala Ala Phe Asn Lys Glu Gly Met Lys Lys  
 100 105 110  
 Asp Asp Gln Ile Ala Ala Ala Met Val Leu Arg Gly Met Ala Lys Asp

115

120

125

Gly Gln Phe Ala Leu Thr Asn Asp Ala Ala Ala His  
 130 135 140

&lt;210&gt; 68

&lt;211&gt; 942

&lt;212&gt; DNA

<213> *Borrelia garinii*

&lt;400&gt; 68

atgagaggat cgcatacaca tcaccatcac ggatccaagg ggactgttaa gaatgctgtt 60  
 gatatgacaa aagctgctgc ggttgctgca agtgctgcaa gtgctgctac tggtaatgca 120  
 gcaattgggg atgttgttaa tggtaatgat ggagcagcaa aagggtggtga tgcggcgagt 180  
 gttaattggga ttgctaaggg gataaagggg attgttgatg ctgctgagaa ggctgatgctg 240  
 aaggaaggga agttgaatgt ggctggtgct gctggtgctg agggtaacga ggctgctggg 300  
 aagctgtttg tgaagaagaa tgctggtgat catggtggtg aagcaggtga tgctgggagg 360  
 gctgctgctg cggttgctgc tgttagtggg gagcagatat taaaagcgat tgttgatgct 420  
 gctaaggatg gtggtgataa gcagggttaag aaggctgagg atgctgaaaa tccgattgac 480  
 gcggtctattg ggagtacggg tgcggatgat aatgctgctg aggcgtttgc tactatgaag 540  
 aaggatgatac agattgctgc tgctatggtt ctgaggggaa tggctaagga tgggcagttt 600  
 gctttgaagg atgctgctca tgataatcat ctgcagccaa gcttaattag ctgagcttgg 660  
 actcctgttg atagatccag taatgacctc agaactccat ctggatttgt tcagaacgct 720  
 cggttgccgc cgggcgtttt ttattggtga gaatccaagc tagcttggcg agattttcag 780  
 gagctaagga agctaaaatg gagaaaaaat cactggatat accaccgttg atatatccca 840  
 atggcatcgt aaagaacatt ttgaggcatt tcagtcagtt gctcaatgta cctataacca 900  
 gaccgttcag ctggatatta cggccttttt aaagaccgta ag 942

&lt;210&gt; 69

&lt;211&gt; 217

&lt;212&gt; PRT

<213> *Borrelia garinii*

&lt;400&gt; 69

Met Arg Gly Ser His His His His His His Gly Ser Lys Gly Thr Val  
 1 5 10 15  
 Lys Asn Ala Val Asp Met Thr Lys Ala Ala Ala Val Ala Ala Ser Ala  
 20 25 30  
 Ala Ser Ala Ala Thr Gly Asn Ala Ala Ile Gly Asp Val Val Asn Gly  
 35 40 45  
 Asn Asp Gly Ala Ala Lys Gly Gly Asp Ala Ala Ser Val Asn Gly Ile  
 50 55 60  
 Ala Lys Gly Ile Lys Gly Ile Val Asp Ala Ala Glu Lys Ala Asp Ala  
 65 70 75 80  
 Lys Glu Gly Lys Leu Asn Val Ala Gly Ala Ala Gly Ala Glu Gly Asn  
 85 90 95  
 Glu Ala Ala Gly Lys Leu Phe Val Lys Lys Asn Ala Gly Asp His Gly  
 100 105 110

Gly Glu Ala Gly Asp Ala Gly Arg Ala Ala Ala Ala Val Ala Ala Val  
115 120 125

Ser Gly Glu Gln Ile Leu Lys Ala Ile Val Asp Ala Ala Lys Asp Gly  
130 135 140

Gly Asp Lys Gln Gly Lys Lys Ala Glu Asp Ala Glu Asn Pro Ile Asp  
145 150 155 160

Ala Ala Ile Gly Ser Thr Gly Ala Asp Asp Asn Ala Ala Glu Ala Phe  
165 170 175

Ala Thr Met Lys Lys Asp Asp Gln Ile Ala Ala Ala Met Val Leu Arg  
180 185 190

Gly Met Ala Lys Asp Gly Gln Phe Ala Leu Lys Asp Ala Ala His Asp  
195 200 205

Asn His Leu Gln Pro Ser Leu Ile Ser  
210 215

<210> 70

<211> 663

<212> DNA

<213> *Borrelia afzelii*

<400> 70

atgagaggat cgcatacacca tcaccatcac ggatccaaga gtgctgtgga tgaggctagc 60  
aagtggtag aagagatgat aacagctgct ggtgaggctg ctacaaaggg tggtagtgg 120  
gaagctagcg aaaagattgg ggatgttggt gataataatc atggtgctgt agctgatgcg 180  
gacagtgtta aggggattgc gaaggggata aaggggattg ttgatgctgc tgggaaggct 240  
tttggtaagg atggtgctgc gaaggatggt gcagctgctg ctggtgatga ggctaacaag 300  
gatgcgggga agttgtttgc tggtcaggat ggtggtggtg ctgatggtga cattgcgaag 360  
gcggctgctg ctgttactgc ggtagtggg gagcagatac tgaaagctat tgttgaggct 420  
gctggtgata aggctaatac ggtgggtgta aaggctgctg gtgcggctac gaatccgatt 480  
gcagctgcga ttgggactga tgatgataat gcggcggcgt ttgataagga tgagatgaag 540  
aagagtaatg ataagattgc tgcggctatt gttttgaggg ggggtggctaa ggatggaaag 600  
tttctgctga atgctaataa taatagtaag gcgagtgtgc tgcagccaag cttaattagc 660  
tga 663

<210> 71

<211> 220

<212> PRT

<213> *Borrelia afzelii*

<400> 71

Met Arg Gly Ser His His His His His His Gly Ser Lys Ser Ala Val  
1 5 10 15

Asp Glu Ala Ser Lys Trp Leu Glu Glu Met Ile Thr Ala Ala Gly Glu  
20 25 30

Ala Ala Thr Lys Gly Gly Thr Gly Glu Ala Ser Glu Lys Ile Gly Asp  
35 40 45

Val Gly Asp Asn Asn His Gly Ala Val Ala Asp Ala Asp Ser Val Lys  
 50 55 60  
 Gly Ile Ala Lys Gly Ile Lys Gly Ile Val Asp Ala Ala Gly Lys Ala  
 65 70 75 80  
 Phe Gly Lys Asp Gly Ala Leu Lys Asp Val Ala Ala Ala Ala Gly Asp  
 85 90 95  
 Glu Ala Asn Lys Asp Ala Gly Lys Leu Phe Ala Gly Gln Asp Gly Gly  
 100 105 110  
 Gly Ala Asp Gly Asp Ile Ala Lys Ala Ala Ala Ala Val Thr Ala Val  
 115 120 125  
 Ser Gly Glu Gln Ile Leu Lys Ala Ile Val Glu Ala Ala Gly Asp Lys  
 130 135 140  
 Ala Asn Gln Val Gly Val Lys Ala Ala Gly Ala Ala Thr Asn Pro Ile  
 145 150 155 160  
 Ala Ala Ala Ile Gly Thr Asp Asp Asp Asn Ala Ala Ala Phe Asp Lys  
 165 170 175  
 Asp Glu Met Lys Lys Ser Asn Asp Lys Ile Ala Ala Ala Ile Val Leu  
 180 185 190  
 Arg Gly Val Ala Lys Asp Gly Lys Phe Ala Ala Asn Ala Asn Asp Asn  
 195 200 205  
 Ser Lys Ala Ser Val Leu Gln Pro Ser Leu Ile Ser  
 210 215 220

<210> 72

<211> 26

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic  
Primer

<400> 72

cggaattcac tcgccttact attatc

26

<210> 73

<211> 29

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic  
Primer

<400> 73

cgggatccga gagtgctggt gatgaggtt 29

<210> 74

<211> 35

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic  
Primer

<400> 74

cgggatccaa gagtgctgtg gatgaggcta gcaag 35

<210> 75

<211> 35

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic  
Primer

<400> 75

ttctgcagca cactcgcctt actattatca ttagc 35

<210> 76

<211> 26

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic  
Primer

<400> 76

cgggatccgc tggtgggagt ygcaac 26

<210> 77

<211> 30

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic  
Primer

<400> 77

aactgcagat tatcatgagc agcatccttc 30

<210> 78

<211> 33

<212> DNA  
 <213> Artificial Sequence  
  
 <220>  
 <223> Description of Artificial Sequence: Synthetic  
         Primer  
  
 <400> 78  
 cgggatccaa ggggactgtt aagaatgctg ttg 33

<210> 79  
 <211> 34  
 <212> DNA  
 <213> Artificial Sequence  
  
 <220>  
 <223> Description of Artificial Sequence: Synthetic  
         Primer  
  
 <400> 79  
 ttctgcagat gattatcatg agcagcatcc ttca 34

<210> 80  
 <211> 17  
 <212> DNA  
 <213> *Borrelia burgdorferi*  
  
 <400> 80  
 tgaggggggct attaagg 17

<210> 81  
 <211> 12  
 <212> DNA  
 <213> Artificial Sequence  
  
 <220>  
 <223> Description of Artificial Sequence: Synthetic  
         Primer  
  
 <400> 81  
 ccggaattcc gg 12